

SEQUENCE LISTING

<110> LEVINE, et al.

<120> VARIANTS Of PROTEIN KINASES

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<150> 09/724,676

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<151> 2000-06-15

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<151> 2000-04-12

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<170> PatentIn version 3.0

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<210> 12
 <211> 750
 <212> DNA
 <213> Homo sapiens

 <220>
 <221> -
 <222> (1)..(750)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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 <211> 794
 <212> DNA
 <213> Homo sapiens

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<210> 14
 <211> 2083
 <212> DNA
 <213> Homo sapiens

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 <211> 1215
 <212> DNA
 <213> Homo sapiens

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ttccaattct	gcaaggcttt taaaattcac cttacatctt ttcaaagcaa gaaaatggaa 180
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cccaaaaatc	ttctgagtgt catctcagga ctttggttat actcatggca cgatggccaa 300
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cacggtggac ggagagagca caagcggaac tgaagacata aagattcagt tcagcaggtc	660
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 <211> 1327
 <212> DNA
 <213> Homo sapiens

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cactgtaatg gttcgtttta atttttcatt ttggattttt ttttaatttc ctttgctact	180
tagattagaa agaacattga tctttcaaac atagatctga atatgaaaga gaaaagaaac	240
acttccatat ttgggttagaa gtctatgtgt ggacagagat ggatacataa tttcacatgc	300
ttggcattcc tttttcacac tttaaaatca ggtaataagt cagtccatct gaggaaagcc	360
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tctgtaataa agtacagcaa aaatactact cgtaagcagt ggctcaaaga gaccctgac	540
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gacagtgttg atggaggtca cgattctgtc attttggatc cagagcgact tgagcctggg	720
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<210> 17
 <211> 364
 <212> DNA
 <213> Homo sapiens

<400> 17	
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ccctgatccc tcagcgttca tgcagcctct tgtccacgga ggctgggtgcc ctgcatgtgc	180
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tccgacctgt gggcgccaat ggccctcccc tgacctcagg gttccttggg ggctgggctg	300
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aaac	364

<210> 18
 <211> 923
 <212> DNA
 <213> Homo sapiens

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gggggtccagg ctctgctgtg tgcgtgcacc agctgcctcc aggccaacta cacgtgtgag 180
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<210> 19
<211> 1739
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(1739)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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acaaagacac aggtcactca aataggcaga gtgacgtcag aatcaagttc gagcacaacg 240
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actctgaggt gtccagacag gtgcggatca aggcttccca gtccgcaggg gatataaata	540
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<210> 20

<211> 1832

<212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(1832)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 20

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ggcactgggg	cttgctccag	attgtggtgg	gagaatgctc	tactaagaga	ttgatggggg	1680
gctgggggtg	agggggggaa	gcctgnagcc	caagagaccc	tgcttctggn	agaatgaatg	1740
gggaatattc	ataaataatg	tacacaaagt	aactctttcc	ttctgctctc	cctgttagct	1800

cccaagtgcc ccccatcaaa cctgggcgcc cg

1832

<210> 21
<211> 1269
<212> DNA
<213> Homo sapiens

<400> 21
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gcagcgactg ggggcggcgc cggcgcgctg gaggcggcca tggcaaagca gtacgactcg 120
gtggagtgcc ctttttgtga tgaagtttcc aaatacgaga agctcgccaa gatcggccaa 180
ggcaccttcg gggaggtggt caaggccagg caccgcaaga ccggccagaa ggtggctctg 240
aagaaggtgc tgatggaaaa cgagaaggag gggttcccca ttacagcctt gcgggagatc 300
aagatccttc agcttctaaa acacgagaat gtggtcaact tgattgagat ttgtcgaacc 360
aaagcttccc cctataaccg ctgcaagggt agtatatacc tgggtgttgc cttctgcgag 420
catgaccttg ctgggctggt gagcaatggt ttggtcaagt tcacgctgtc tgagatcaag 480
agggtgatgc agatgctgct taacggcctc tactacaacc acgacttctt ctggtccgac 540
cccatgccct ccgacctcaa gggcatgctc tccaccacc tgacgtccat gttcgagtac 600
ttggcaccac cgcgcgggaa gggcagccag atcaccacc agtccaccaa ccagagtcgc 660
aatcccgcca ccaccaacca gacggagttt gagcgcgtct tctgagggcc ggcgcttgcc 720
actagggctc ttgtgttttt tttcttctgc tatgtgactt gcacgtgga gacagggcat 780
ttgagtttat atctctcatg catattttat ttaatcccca cctggggctc tgggagcagc 840
ccgctgagtg gactggagtg gagcattggc tgagagacca ggagggcact ggagctgtct 900
tgtccttgct ggttttctgg atggttccca gagggtttcc atggggtagg aggatgggct 960
cgcccaccag tgactttttc taagagctcc cggcgtggtg gaagagggga caggtccttc 1020
accacccac aatcctattc tcgggctgag aaccctgcgt ggggacaggg ctgcctcag 1080
gaatgggctg tttttggcct aaccctcaga aacactgggg ctggcacaaa ctcttggttt 1140
cttcaacagg agaattttac tgtgtttctt ttggttccat tgtttgagga cattcctggg 1200
cacagtttgg tccgttagaa ttaaaagttg aatttttttt ttttttwaat tttttttttt 1260
ycccccaaa 1269

<210> 22
<211> 623

<212> DNA
 <213> Homo sapiens

 <220>
 <221> -
 <222> (1)..(623)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 22
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 gggggcctct ctagcttgcg gcctgtgtct atggtcgggc cctctgcgtc cagctgctcc 120
 ggaccgagct cgggtgtatg gggccgtagg aaccggctcc ggggccccga taacgggccg 180
 cccccacagc accccgggct ggcgtgaggg tctcccttga tctgagaatg gctacctctc 240
 gatatgagcc agtggctgaa attggtgtcg gtgcctatgg gacagtgtac aaggcccgtg 300
 atccccacag tggccacttt gtgccctcaa gagtgtgaga gtccccaatg gaggaggagg 360
 tggaggaggc cttcccatca gcacagttcg tgagggtggct ttactgaggc gactggaggc 420
 ttttgagcat cccaatgttg tccggctgat ggacgtctgt gccacatccc gaactgaccg 480
 ggagatcaag gtaaccctgg tgtttgagca tgtagaccag gacctaagga catatctgga 540
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 ttganagggtg gattgggacc ttt 623

<210> 23
 <211> 502
 <212> DNA
 <213> Homo sapiens

<400> 23
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 ggaccgagct cgggtgtatg gggccgtagg aaccggctcc ggggccccga taacgggccg 180
 cccccacagc accccgggct ggcgtgaggg tctcccttga tctgagaatg gctacctctc 240
 gatatgagcc agtggctgaa attggtgtcg gtgcctatgg gacagtgtac aaggcccgtg 300
 atccccacag tggccacttt gtgccctcaa gagtgtgaga gtccccaccc acctctcctt 360
 ttgaggcttc tcttctcct tccatttct ctacactaag gggatatgttc cctcttgtcc 420
 ctttccctac ctttatattht ggggtccttt tttatacagg aaaaacaaaa caaagaaata 480
 aagtcgacgc ggccgcgaat tc 502

<210> 24
 <211> 1148
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(1148)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 24
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 cctgggaaga ctcccttctg ctccccaaaa cccaaggcc tggctcgggg ccactggagc 180
 cgcaggcggg acatatgtgt gaccggccct ctgcccctgg cagccccgcg ctgtgtactg 240
 taaggacgtg ctggacatcg agcagttctc cactgtgaag ggcgtcaatc tggaccacac 300
 agacgacgac ttctactcca agttctccac gggctctgtg tccatcccat ggcaaaacga 360
 gatgatagaa acagaatgct ttaaggagct gaacgtgttt ggacctaatg gtaccctccc 420
 gccagatctg aacagaaacc accctccgga accgcccag aaagggtgtc tccagagact 480
 cttcaagcgg cagcatcaga acaattccaa gagttcgccc agctccaaga ccagttttaa 540
 ccaccacata aactcaaacc atgtcagctc gaactccacc ggaagcagct agtttcggct 600
 ctggcctcca agtccacagt ggaaccagcc cagacccttc tccttagaag tggaaagtagt 660
 ggagcccctg ctctggtggg gctgccaggg gagaccccgg gagccgggga aggaggccgt 720
 ccatcccgtc gacgtagaac ctcgaggttt ctcaaagaaa tttccactca ggtctgtttt 780
 ccgaggcggc cccggccggg gtggattgga tttgtctttg gtgaacattg caatagaaat 840
 ccaattggat acgacaactt gcacgtatth taatagcgtc ataactagaa ctgaattttg 900
 tctttatgat ttttaaagaa aagttttgta aatttctcta ctgtctcagt ttacattttg 960
 tatatttgta tttaaatgaa agtgagactt tgaggggtgta tattttctgt gcagccactg 1020
 ttaagccatg tgttccaagg catttttagcg gggagggggg tatcaaaaaa aaaaaaatgt 1080
 gactcaagac ttccagagcc tcaaatgaga aaatgtcttt attaaatgta gaaagtgatc 1140
 catacttc 1148

<210> 25
 <211> 1679
 <212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(1679)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 25

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ggcattttcc tgtgtcctcc cacacctgca ctctcccaag gctcttgagg tcactctgag	180
atggcagatt ggggtccctg ttgtccttgg acagatgaga atgccgagag ctcgtatgcc	240
tnggccaagg gcacacagca aggccgggtg cccatgcggc tgtcccaggg acccactgac	300
cctgctgtcc ccctcaggcc acattaggat ctccagacctg ggcttggtctg tgaagatccc	360
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gaacaaccag aggtacggcc tgagccccga ctactggggc cttgggtgcc tcatctatga	480
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gtccatctgc aagatgctgc tcacgaaaga tgcgaagcag aggctgggct gccaggagga	660
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cttctactcc aagttctcca cgggctctgt gtccatccca tggcaaaacg agatgataga	900
aacagaatgc ttttaaggagc tgaacgtgtt tggacctaat ggtaccctcc cgccagatct	960
gaacagaaac caccctccgg aaccgcccaa gaaagggctg ctccagagac tcttcaagcg	1020
gcagcatcag aacaattcca agagtccgcc cagctccaag accagtttta accaccacat	1080
aaactcaaac catgtcagct cgaactccac cggaagcagc tagtttcggc tctggcctcc	1140
aagtccacag tggaaaccag ccagaccctt ctcccttagaa gtggaagtag tggagccctt	1200
gctctgggtg ggctgccagg ggagaccccg ggagccgggg aaggaggccg tccatcccg	1260
cgacgtagaa cctcgagggt tctcaaagaa atttccactc aggtctgttt tccgaggcgg	1320
ccccggccgg ggtggattgg atttgtcttt ggtgaacatt gcaatagaaa tccaattgga	1380
tacgacaact tgcacgtatt ttaatagcgt cataactaga actgaatttt gtctttatga	1440

tttttaaaga aaagttttgt aaattttctct actgtctcag tttacatttt gtatatattgt	1500
atttaaataa aagtgagact ttgaggggtgt atattttctg tgcagccact gttaagccat	1560
gtgttccaag gcatttttagc ggggaggggg ttatcaaaaa aaaaaaaatg tgactcaaga	1620
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<210> 26
 <211> 897
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(897)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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ggcattttcc tgtgtcctcc cacacctgca ctctcccaag gctcttgtgg tcaactctgag	180
atggcagatt ggggtccctg ttgtccttgg acagatgaga atgccgagag ctcgatatgcc	240
tnggccaagg gcacacagca aggcgggtg cccatgcggc tgtcccaggg acccactgac	300
cctgctgtcc ccctcaggcc acattaggat ctgagacctg ggcttggctg tgaagatccc	360
cgagggagac ctgatccgcg gccgggtggg cactgttggc tacatggctc cagaggctct	420
gaacaaccag aggtacggcc tgagccccga ctactggggc cttggctgcc tcatctatga	480
gatgatcgag ggccagtcgc cgttccgcgg ccgcaaggag aagggtgaagc gggaggaggt	540
ggaccgccgg gtcttgagga cggaggaggt gtactccac aagttctccg aggaggccaa	600
gtccatctgc aagatggtga gctcctggtg gccagatgcc accctcaagc tgggtggctcc	660
cctccctggg cctggcccca gtctgtcccc agaacagcaa acaggctgaa gggacagggg	720
tctgagcatg ggggtggggg ttgcagccca ccaagctaga gttgagggca ctcttctgtg	780
aggactgggg tggtcagcgg agacttccag gagaaagcct ggggtggggca gggacaccca	840
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<210> 27
 <211> 1224
 <212> DNA
 <213> Homo sapiens

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 <212> DNA
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2027

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<211> 1995
<212> DNA
<213> Homo sapiens

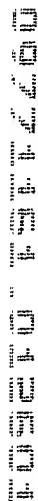
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<222> (1)..(1995)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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<211> 940
<212> DNA
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<212> DNA
<213> Homo sapiens

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<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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<210> 36
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<212> DNA
<213> Homo sapiens

<220>
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<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

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<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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 <212> DNA
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 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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cggccccgcc	gcagaccagc	tggcgggtgt	ggagaccagg	ctcctgaccc	cgccatgcat	1140
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<210> 48
 <211> 1154
 <212> DNA
 <213> Homo sapiens

<400> 48	
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agaccctgca	ccgggcttgg actcgcagcc gggactgacg tgtagaacia tcgtttctgt 180
tggaagaagg	gtttttccct tctttttggg gtttttggtg cttttttttt ttcttttttc 240
tttgtaaaat	tttggagaag ggaagtccga acacaaggaa ggaccgctca cccgcggact 300
cagggctggc	ggcgggactc caggaccctg ggtccagcat ggagggtggtg gaccgcagc 360
agctgggcat	gttcacggag ggcgagctga tgcggtggg tatggacacg ttcattccacc 420
gcattcgactc	caccgaggtc atctaccagc cgcgccgcaa gcgggccaag ctcatcgga 480
agtacctgat	gggggacctg ctgggggaag gctcttacgg caagggtgaag gaggtgctgg 540
actcggagac	gctgtgcagg agggccgtca agatcctcaa gaagaagaag ttgcgaagga 600
tccccaacgg	ggaggccaac gtgaagaagg aaattcaact actgaggagg ttacggcaca 660
aaaatgtcat	ccagctggtg gatgtgttat acaacgaaga gaagcagaaa atgtatatgg 720
tgatggagta	ctgcgtgtgt ggcattgcagg aaatgctgga cagcgtgccg gagaagcgtt 780
tcccagtgtg	ccaggccccac gggtaacttct gtcagctgat tgacggcctg gactacctgc 840
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gcaccctcaa	aatctccgac ctgggcgtgg ccgaggtagg cacgtgctag ggggggccct 960
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<210> 49
 <211> 930
 <212> DNA
 <213> Homo sapiens

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 gccaaaggagg gacagaaatg gatggcttta ggagaacccat agaaaaccag cactctcgta 240
 atgatgtcat ggtttctgag tggctaaaca aactgaatct agaggagcct cccagctctg 300
 ttcctaaaaa atgcccagagc cttaccaaga ggagcagggc acaagaggag caggttccac 360
 aagcctggac agcaggcaca tcttcagatt cgatggccca acctccccag actccagaga 420
 cctcaacttt cagaaaccag atgcccagcc ctacctcaac tggaacacca agtcctggac 480
 cccgaggggaa tcagggggct gagagacaag gcatgaactg gtctctcagg accccggagc 540
 caaatccagt aacaggggcga ccgctcgta acatatacaa ctgctctggg gtgcaagttg 600
 gagacaacaa ctacttgact atgcaacaga caactgcctt gccacatgg ggcttggcac 660
 cttcgggcaa ggggaggggc ttgcagcacc ccccaccagt aggttcgcaa gaaggcccta 720
 aagatcctga agcctggagc aggccacagg gttggtataa tcatagcggg aaataaagca 780
 ccttccaagc ttgcctccaa gagttacgag ttaaggaaga gtgccacccc ttgaggcccc 840
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 gagtcaataa acatgatgga atgctaaaaa 930

<210> 50
 <211> 2616
 <212> DNA
 <213> Homo sapiens

<400> 50
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 gactgcaggc gcatgccacc atggctcagct aatttaaaaa aaaaatgttt ttggctgggc 180

aaagtggatc acatctgtaa tcccagcacc tagggatgcc aaggcaagaa gattgcttgt	240
gagcccagaa gttcgagacc agcctgggca acatggtgaa actctacctc taccaaaaaa	300
atgtaaaaat tagccatatt tgacctcaag tctagacaga acttctttgt atattttaga	360
gaggagcaag caagggagct gtacaggaga ctaagggaaa aacctcgaga ccagcgaact	420
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aaagtgcgag tgatctatac gcagctcagt aaaactgtgg ttgcaagca gaaggcgctg	540
gaactgttgc ccaagggtga agagggtgtg agcttaatga atgaggatga gaagactgtt	600
gtccggctgc aggagaagcg gcagaaggag ctctggaatc tcctgaagat tgcttgtagc	660
aagggtccgtg gtcctgtcag tggaagcccg gatagcatga atgcctctcg acttagccag	720
cctgggcagc tgatgtctca gccctccacg gcctccaaca gcttacctga gccagccaag	780
aagagtgaag aactgggtggc tgaagcacat aacctctgca ccctgctaga aaatgccata	840
caggacactg tgagggaaaca agaccagagt ttcacggccc tagactggag ctggttacag	900
acggaagaag aagagcacag ctgcctggag caggcctcat gatgtggggg gactcgaccc	960
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gttcctgctg cactgatggc ccaggggtct ctggtatcca gatggagctc tcgcttcctc	1140
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cggtttgcca cacacgtgac tggacagtgt ccaattcaaa tctttcaggg cagagtccga	1380
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caatccactg ttagaatacc tatgggttagg gcttctgaac taaaataatg gaaaatttta	1980
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acaatttttg aaaccctggg aaatgacagt gggaaataac acccgaaagg caaggacggg	2460
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<210> 51
 <211> 1323
 <212> DNA
 <213> Homo sapiens

<400> 51	
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gccagaggat ctatgatggc cagtacctcc aggccctgaa cgcggactgg cacgcagact	240
gcttcagggtg ttgtgactgc agtgccctcc tgtcgcacca gtactatgag aaggatgggc	300
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ggacccctcc atcctccttc cttcccagtc tatggaaaca cagtgggaagg ggtatctggc	1260
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<210> 52
 <211> 1262
 <212> DNA
 <213> Homo sapiens

<400> 52	
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cctggagggga agaacgtatg ggagaggaag gaagcgagtt gcccggtgtg gcaagctgcg	180
gccagaggat ctatgatggc cagtacctcc aggccctgaa cgcggactgg cacgcagact	240
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agcaaatac caagggactg gttatgggtg ctggggagct gaagtaccac cccgagtgtt	420
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cggtctgtgg caccgagcac tcacacaccg tccgcgtcca gggagtggat ccgggctgca	720
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gaattccggc	ccagtgggga	gctgttcgcc	atcaaggctc	tgaagaaagg	ggacattgtg	2040
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gcgggacacc	ccttctctgt	gaacctcttc	ggctgtttcc	agacaccgga	gcacgtgtgc	2160
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ttcgacttcg	tggccggggg	ctgctagccc	cctcccctgc	cctgcccct	gccctgccc	2460
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<210> 54
 <211> 1464
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -

<222> (1)..(1464)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 54
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 tggccccctgc tagagcagct gggcctggcc ggggcagacc tggcggcccc cggggtacag 180
 cagcagctgg agctggagcg ggagcggctg cggcgggaaa tccgcaagga gctgaagctg 240
 aaggagggtg ctgagaacct gcggcggggc accactgacc tgggcccgcag cctgggcccc 300
 gtagagctgc tgctgcgggg ctccctgcgc cgccctgacc tgctgcacca gcagctgcag 360
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 agtaagacca agattgacat catccgcatg caactccgcc gggcgctgca ggccgaccag 660
 ctggagaacc aggcagcccc ggatgacacc caaggagtc ctgacctggg ggctgtggag 720
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 gctgcggcct cctccgctgc cttcagcacc cgccctggcc ggccctttcc cgccacgcac 1020
 tacagcacc cgtgcaagcc cgcgcgcctc acagggaccc tggaggtacg agtgggtggc 1080
 tgcagagacc tcccagagac catcccgctg aaccctaccc cctcaatggg gggacctggg 1140
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 ggaagcctca gtggccggag cagcctcaaa gcagaagccg agaacaccag tgaagtcagc 1260
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 ccgcggagca ggcagccttc ctggacttcg acttcgtggc cgggggctgc tagccccctc 1380
 cctgccccct gcccctgccc ctgcccgaga gctcttagtt tttaaaaagg cctttgggat 1440
 ttgccggatc cttgcaccc caaa 1464

<210> 55

<211> 1080
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(1080)
<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

<400> 55
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gggtgtgagg tagctccagg tccaaaaggt cacaagaggt catgttctcc agctgtggga 180
gccccaggag agaggcaagg cgnncttttg gtcaagcaa agcttctctg tggagggcag 240
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gagaagtgca gagagtaagg tgccttatg ttggaaactc aagtggaagg aagatttggg 480
ttggttttat tctcagagcc attaaacact agttcagtat gtgagatata gattctaaaa 540
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atggaaggaa gtggaaccag atgcagaaga ggaaatgatg gaaggactta tggatcaga 1020
taccaatatt taaaagtttg tataataata aagagtatga ttgtggttca aggataaaaa 1080

<210> 56
<211> 1665
<212> DNA
<213> Homo sapiens

<400> 56
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ggatcgttgc aggaaggttc taagggctca ggagtgtgag tggcccaggc acctgccgtc 180
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 agcccaggac cccactgagc cccacgcct gtctcgcagc ctgagcgggtg catccccgtt 300
 cctgggcgag accaagcagg agacgctcac caacatctca gccgtgaact acgacttcga 360
 cgaggagtac ttcagcaaca ccagcgagct ggccaaggac ttcattcgcc ggctgctcgt 420
 caaagatccc aagcggagaa tgaccattgc ccagagcctg gaacattcct ggattaaggc 480
 gatccggcgg cggaacgtgc gtggtgagga cagcggccgc aagcccgagc ggcggcgcct 540
 gaagaccacg cgtctgaagg agtacaccat caagtcgcac tccagcttgc cgcccaacaa 600
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 cgccatctac gaggagaagg aggcctggta ccgcgaggag agcgacagcc tgggccagga 780
 cctgcggagg ctacggcagg agctgctcaa gaccgaggcg ctcaagcggc aggcgcagga 840
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 ttggtgggggt cttccactgt gtgcccttct cgccgaggcc ggtcccccg gtgtgggggtg 1560
 cctgctgcg gactcctccg cgagcccat cgtcgcgcct gtggacgcct aggcaagagc 1620
 ggccctctgc agccaagaga aataaaatac tggcttcag ataaa 1665

<210> 57
 <211> 2081
 <212> DNA

<213> Homo sapiens

<400> 57

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tccccctccc tacctcccca gaatgtaccc tacagccaaa gtctcagcag cctacaggga	180
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tgcttgccag cggaaccggg ctgtgtgcca gcttctggtg gatgcaggag catctctgag	360
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aactgctgtt tgaccctggt attcgggcaa agaggacatg agcaagcgta tcacatctgc	540
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ctcctgcaag attctacctg agaccatgcc actagctttt aagggtacc aagatgtaca	660
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acgagttcca caaaatttga tccttattgc ttccagcaag tagcatgaac ttctgtgttc	780
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tccccctatg cattgggtttt ttccctgta ccatacaatt ctactgtaac taccatcaa	900
cttaaagaaa aatattatct cttctcttta cattcagtct tggaagacca caagattgtc	960
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gccaacctc cgtttcatta tataggagct ggggaagtgcc acatggataa tgtcaacttg	1140
tgtgctatat ctctgaggaa tggtaggtg gcatgggaga tgtctgtgct tggaggtagc	1200
tcagagaggt aaccagggg tcagcccagg ctgctgggct gtagccaata gccatgcagg	1260
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ttttgctgca atagaagatg agcaaaggat taaacagagg cccacagcta gtttgagaa	1380
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aagcaccta attgacttgg aaaaaaaaaa cagcaaaagc aaaagtagca acatatgtca	1560
acatatgtca ctgaaatagg aaacagtcac tggaatgttg cacagaggct aatagctatg	1620

ccaaaggaaa aataagatga cctaatactact gtctctgggtt tctctgaact tattttttatt 1080
 ttttaaaaat taagggtc 1097

<210> 59
 <211> 3382
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(3382)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 59
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 gtggggcggc gcggctgggg cgccggggcc ggagttgcct cccggggccc gcgctgaggg 180
 ccccgccgcy gccgcccgtg ctgcttctgc ttgcgctgtt gccgctgctg cccgcgcctg 240
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 gcgtgagtct ctacctgagc gaggacgagg tgcgccggct gatcggctctt gatgcagaac 360
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 ccagtgagac aaatttctg cacttcacct ggcatgcgaa gtccaagggtt gaataaagc 480
 tgggattcca agtggaacaat gttttggcaa tggatatgcc ccaggtcaac atttctgttc 540
 agggggaagt tccacgcact ttatcagtgt ttcgggtaga gctttcctgt actggcaaag 600
 tagattctga agttatgata ctaatgcagc tcaacttgac agtaaatct tcaaaaaatt 660
 ttaccgtctt aaattttaaa cgaaggaaaa tgtgctacaa aaaacttgaa gaagtaaaaa 720
 cttcagcctt ggacaaaaac actagcagaa ctatttatga tcctgtacat gcagctccaa 780
 ccacttctac gcgtgtgttt tatattagtg taggggtttg ttgtgcagta atatttctcg 840
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 atctgagagc agacacgccc aacaatgcaa ctctatcac cagctcctag ttatcctacc 1020
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 aaggatatag caatatccag agagaggata actctaaaag atgtactcca agaagggtact 1140
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gcatttgtca aaacagttaa agatcaagct tctgaaattc aggtgacaat gatgctcact 1260

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tctaacacaa cgccaacaga agcacatttg tcttccagaa caccgtgcct tagaaatgct 1920

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gacacagata tttggaatta gctatcttag ggccaactgc tttttatttt ttacttcat 2160

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atattagagg aagaaaattt aaagaaaagc tagaggaaaa aaaaattttt ttaaaaatac 2460

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aatgtgtagc agttngaggc ttttgctggt tttaaaaaag ccttatgaat cagcagcaca	3300
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ctgtggtaca ttttatttaa ca	3382

<210> 60
 <211> 2195
 <212> DNA
 <213> Homo sapiens

<400> 60	
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tgtgtgttgc tggggatctg agaagtcgat ctttgagctg agcgctggtg aaggagaaac	120
aagccatgga aggaaagggtg ccaagtgggtc aggcgagagc ctccaggga aaggccttgg	180
gcagggtggga atcctgattt gttcctgaaa ggtagtttgg ctgaatcatt cctgagaagg	240
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<210> 62
<211> 1149
<212> DNA
<213> Homo sapiens

<400> 62
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ctgcagccga gacctggccc ggcgcggct cctgcagca tgcccagcct ccgcccgcgc	300
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 <211> 765
 <212> DNA
 <213> Homo sapiens

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ttctccaagt tgggggctca gaggggagtc atcatgagcg atgttaccat tgtgaaagaa	180
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<210> 65
<211> 968
<212> DNA
<213> Homo sapiens
<220>
<221> -
<222> (1)..(968)
<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

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ttctccaagt tgggggctca gaggggagtc atcatgagcg atgttaccat tgtgaaagaa 180
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ccctcaaca acttttcagt ggcaaaatgc cagttaatga aaacagaacg accaaagcca 360
aacacattta taatcagatg tctccagtgg actactgtta tagagagaac atttcatgta 420
gatactccag aggaaaggga agaattggaca gaagctatcc aggctgtagc agacagactg 480
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agtggaaaat actatgctat gaagattctg aagaaagaag tcattattgc aaaggtaact 720
gatttattaa agttgattac taaatttttg tttgcagtgt gcatgtgttt gtgggctcat 780

gaatttacat gctaattgtat gcaaattcca ttaaacaacc naaatatggt tgnagactac	840
tgctacagta atttttgtgt attaataattt gtaattttta aagttttcag acattcataa	900
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tgtttggg	968

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 <211> 2410
 <212> DNA
 <213> Homo sapiens

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ttttatgttt gtgttaattt cccccactaa atcagtaatt attacaatcc tgtccctgct	300
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<210> 67
 <211> 798
 <212> DNA
 <213> Homo sapiens

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 tccgccccct tccgcctcc ccgtatataa gacttcgccg agcactctca ctgcacaag 180
 tggaccgggg tgttgggtgc tagtcggcac cagaggcaag ggtgcgagga ccacggccgg 240
 ctcgacgtg tgaccgcgcc tagggggtg cagcgggcag tgcggggcgg caaggcgacc 300

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gaatcgagc cacctcagtc ccaggcgcaa gtgcccccg cgccccctca ccaccatcac	480
caccattcgc actcggggcc ggagatctcg cggattatcg tcgacccac gactgggaag	540
cgctactgcc ggggcaaagt gctgggaaag ggtggctttg caaaatgtta cgagatgaca	600
gatttgacaa ataacaaagt ctacgccgca aaaattattc ctcacagcag agtagctaaa	660
cctcatcaaa gggaaaaggt gtgtatgact cttgagtaaa gtattttctt tgtgtgcaag	720
gatggccctt ccctgttagg aaaatgtctt ctgcatgtgt aatcactggc ttttccgagg	780
tgactggaag ctaataag	798

<210> 68
 <211> 877
 <212> DNA
 <213> Homo sapiens

<400> 68	
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tccgccccct tccgcctcc ccgtatataa gacttcgccg agcactctca ctgcacaaag	180
tggaccgggg tgttgggtgc tagtcggcac cagaggcaag ggtgcgagga ccacggccgg	240
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cgctactgcc ggggcaaagt gctgggaaag ggtggctttg caaaatgtta cgagatgaca	600
gatttgacaa ataacaaagt ctacgccgca aaaattattc ctcacagcag agtagctaaa	660
cctcatcaaa gggaaaagat tgacaaagaa atagagcttc acagaattct tcatcataag	720
catgtagtgc agttttacca ctacttcgag gacaaagaaa acatttacat tctcttgga	780
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<210> 69
 <211> 1349
 <212> DNA
 <213> Homo sapiens

<400> 69
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 tccgccccct tccgcctcc ccgtatataa gacttcgccg agcactctca ctcgcacaaag 180
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 agatacttgt ggctggaaaa gtgcattcct tgtaataaaa cttttttattt attacagccc 1260
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 gcaataaaga gtatgaaaac gcagaaaaa 1349

<210> 70
 <211> 538
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(538)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 70
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 gctgctgcta ccattcttca aggtcagaaa gactcctgtt tatggagggg gttgcagcag 480
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<210> 71
 <211> 3308
 <212> DNA
 <213> Homo sapiens

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<210> 72
 <211> 3503
 <212> DNA
 <213> Homo sapiens

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<210> 73
<211> 2544
<212> DNA
<213> Homo sapiens

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<220>
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<222> (1)..(2544)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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 <211> 2324
 <212> DNA
 <213> Homo sapiens

 <220>
 <221> -
 <222> (1)..(2324)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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<210> 75
 <211> 1396
 <212> DNA
 <213> Homo sapiens

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1396

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<211> 513
<212> DNA
<213> Homo sapiens

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<210> 77
<211> 2044
<212> DNA
<213> Homo sapiens

<400> 77
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 <211> 934
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 <213> Homo sapiens

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agattttaccc cccaaaaaaa attgtcaatg agaaataaag ctaactgata tcaaaaagca	180
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<210> 79
 <211> 1032
 <212> DNA
 <213> Homo sapiens

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ctaggggaca taaaattagt gattactttg agacagcccc tctatggttt agatggcagt	420
gctgcaaagg aggcaacaga ggagcagtct gctctgcaa tcctcacgct agtgatgcta	480
gcaaaacctc ggcttgacac agagcagctg gcgtaaagga gagctggcct ctgcttcact	540
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gaaaaaatat ctgcactaga aaacagtaag aattctgact tagagaagaa ggaggggaaga	720
atagatgatt tattaagagc caactgtgat ttgagacggc agattgatga acagcaaaaag	780
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atagaaaagg ttagtgaata atgttgggtct aaactctgta tccaagata ctcaatgtgt	900
gtcattgtgt ggcttcttat tccttacttg agatgaaaat atttaaaagt agagcttttt	960
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<210> 80
 <211> 2234
 <212> DNA
 <213> Homo sapiens

<400> 80	
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gcagatgcct gggctgctcc catcgccatg cagatctaca agaagcacct ggaccccagg	180
cccgggccct gccacctgag ctgggcctgg gcctgggcca gctggcctgc tgctgcctgc	240
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<210> 81
 <211> 2608
 <212> DNA
 <213> Homo sapiens

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tacagcccgg gcaattcagt ttctacatca ggacagcccc agcctcatcc atggagacat	180
caagagttcc aacgtccttc tggatgagag gctgacaccc aagctgggag actttggcct	240
ggcccgggtc agccgctttg ccgggtccag cccagccag agcagcatgg tggcccggac	300

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catgatgaga	ccctgtctct	gccaaaaaat	tttttaaact	attagcctgg	cgtggtagcg	1920
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gtccaaaagt agcccagggc tgtagcacag gcttcacagt gattttgtgt tcagccgtga	2520
gtcacactac atgccccctg gaagctgggc attggtgacg tccaggttgt ccttgagtaa	2580
taaaaacgta tgttgcaatc tcgggaaa	2608

<210> 82
 <211> 1237
 <212> DNA
 <213> Homo sapiens

<400> 82	
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cgcccagcac ttcttgtaag aggtgccgcc ctgggtcatg tgccgcttct acaaagtgat	180
ggacgccttg gagcccccg actggtgcca gtgcgccgcc ctgatcgtgc gcgaccagac	240
cgagctgcgg ctgtgcgagc gctccgggca gcgcacggcc agcgtcctgt ggccctggat	300
caaccgcaac gcccggtgtg ccgacctcgt gcacatcctc acgcacctgc agctgctccg	360
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gttgccatcc tcagcctcca ccttcctctc ccagctttt ccaggctccc agaccattc	540
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ggtgtatgct gtgaagaggc tgaaggagaa cgctgacctg gagggtgact cagtgaagca	840

gagcttcctg accgaggtgg agcagctgtc caggtttcgt cacccaaaca ttgtggactt	900
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ctcttgctg tccaaaagta gccccagggt gtagcacagg cttcacagtg attttgtgtt	1140
cagccgtgag tcacactaca tgccccctg aagctgggca ttggtgacgt ccagggtgtc	1200
cttgagtaat aaaaacgtat gttgcaatct cgggaaa	1237

<210> 83
 <211> 1286
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(1286)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 83	
gaggggacct taaaaattac cggccacaaa aagaaaataa atttaggaaa aattattggt	60
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tgcttttagac catctacata gcttggaat aatttataga gacttaaaac cagaaaatat	180
acttcttgat gaagaaggtc acatcaagtt aacagatttc ggcctaagta aagagtctat	240
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agttaatcgt cgaggtcata ctacagagtgc tgactgggtg tcttttggtg tgtaaatgtt	360
tgaaatgctt actggtacac tccctttcca aggaaaagat cgaaaagaaa caatgactat	420
gattcttaaa gccaaacttg gaatgccaca gtttttgagt cctgaagcgc agagtctttt	480
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agaaattaaa agacattcat ttttctcaac gatagactgg aataaactgt atagaagaga	600
aattcatccg ccatttaaac ctgcaacggg caggcctgaa gatacattct attttgatcc	660
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tcagcttttt cgggggttta gttttgttgc tattacctca gatgatgaaa gccaaagctat	780
gcagacagtt ggtgtacatt caattgttca gcagttacac aggaacagta ttcagtttac	840
tgatggatat gaagtaaaag aagatattgg agttggctcc tactctgttt gcaagagatg	900

tatacataaa gctacaaaca tggagtttgc agtgaaggta aatTTTTTTT atttaaaatg	960
caattcatac agttcttggt catgcatgtc agtaccagtt aaaaattaca ctcccccttgt	1020
tgttaaaagt gccttttggt ataaaaaagt taaatatctg gctagtgatc ttcagagatc	1080
ttaatctaga accctgtgag ctaaaggtaa ggtgggtata tatctagttt tcccagagca	1140
gtagcagttt acacctcaag tgatTTTTTT tctTTTTTTT cctcaagtga tttttaaaagt	1200
atctttttac tctgagaagt ccccatTTTT tgctcanggt gtcagcaaAT tcctcaaaat	1260
tgtgtgcaaa attttgtatg tttaaa	1286

<210> 84
 <211> 752
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(752)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 84	
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agcgcgtgaga atggacagca aattatggat gaacctatgg gagaggagga gattaaccca	120
caaactgaag aagtcagtat caaagaaatt gcaatcacac atcatgtaaa ggaaggacat	180
gaaaaggcag atccttccca gtttgaactt ttaaaagtat tagggcaggg atcatttggga	240
aagggtttct tagttaaaaa aatctcaggc tctgatgcta ggcagcttta tgccatgaag	300
gtattgaaga aggccacact gaaagttoga gaccgagttc ggacaaaaat ggaacgtgat	360
atcttggtag aggttaatca tccttttatt gtcaagttgc attatgcttt tcaaactgaa	420
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atcatgctac taagttgaat acaatgtaat atgattgttt aggagattat aaaaaatcaa	660
gtggcttcat gaaactccca cagtaatgtn tagcgtgcct gtgcttcaca tctctgctaa	720
cactgtagtt tcatacttta aatnactcag tt	752

<210> 85
 <211> 1826
 <212> DNA

<213> Homo sapiens

<400> 85

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ccaagaaaag gaagctccgt cccttcccgc tcaccggct tccccacccc ttgtactcta	120
aactctgcas agggcgagcg ygcggccack gakgcgccga ggaggagcga gcgccgccgg	180
gcagcggcgt gccctcgggg gagagggcgc cggakargag cggcggcgcg gcggcgakgg	240
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ggccaggcgg tgcggctggg cgggggacgc cgccgccgtt gctgcccggc ccggagagat	360
gagcacggag gcggacgarg gcatcacttt ctctgtgcc ccttcgccc cctcgggctt	420
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ccaagggcaa ctggtggtgg ccgagagcga ggccctgcag agcttgcggg aggcgtgcga	720
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catcctctac tgygatacta actcggactc tctgcagtca ctgaaggaaa tmatttgcca	960
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aacctttgat ttggcctccc atcaccatgt gaagtttcat tatgcatttg cactgaatag	1440
gagaaatctc cctggtgaca gagcaaaagc tcttgatatt atgattccca tggtgcaaag	1500
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ggcatttgaa tctgagccaa cactacagtc aggaattaat tatgcggtcc tcctcctggc 1680
agctggacac cagtttgaat ctcccttga gctccggaaa gttggtaatt acaacttgat 1740
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attctgacgg ctctccaggt tttgtc 1826

<210> 86
<211> 476
<212> DNA
<213> Homo sapiens

<400> 86
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agatgcagag cacttctaata catctgtggc ttttatctga tatttttaggc caaggagcta 120
ctgcaaagt ctctcgtgga agacataagt ggatgttcaa atgagagaat ttgaagtgtt 180
gaaaaaactc aatcacaaaa atattgtcaa attatttgct attgaagagg agacaacaac 240
aagacataaa gtacttatta tggaattttg tccatgtggg agtttataca ctgttttaga 300
agaaccttct aatgcctatg gactaccaga atctgaattc ttaattgttt tgcgagatgt 360
ggcgggtgga atgaatcatc tacgagagaa tggatatagt caccgtgata tcaagccagg 420
aaatatcatg cgtgcactat accattctct cgtagatgat tcattccacc caccac 476

<210> 87
<211> 2131
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(2131)
<223> "n" can be any nucleotide 'a', 'c', 'g' or 't'

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caggatctta cttggatcca gagtgctagc tgcaaaagct tctgggaatt gtacacttaa 180
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gtggccatta agaagctcag cagacccttt cagaaccaa cacatgccaa gagagcgtac 300
cgggagctgg tcctcatgaa gtgtgtgaac cataaaaaa ttattagttt attaaatgtc 360
ttcacacccc agaaaacgct ggaggagttc caagatgttt acttagtaat ggaactgatg 420

gatgccaaact tatgtcaagt gattcagatg gaattagacc atgagcgaat gtcttacctg	480
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ctggccagga cagcaggcac aagcttcatg atgactccat atgtggtgac acgttattac	660
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gaccagtggga ataaggtaat tgaacaacta ggaacaccat gtccagaatt catgaagaaa	840
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cccaaactct tcccagattc cctcttccca gcggactccg agcacaataa actcaaagcc	960
agccaagcca gggacttgtt gtcaaagatg ctagtgattg acccagcaaa aagaatatca	1020
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<212> PRT
<213> Homosapiens

<400> 92

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Leu	Phe 35	Val	Ser	Thr	Leu	Asp	Gly 40	Ser	Leu	His	Ala	Val 45	Ser	Lys	Arg
Thr	Gly 50	Ser	Ile	Lys	Trp	Thr 55	Leu	Lys	Glu	Asp	Pro 60	Val	Leu	Gln	Val
Pro 65	Thr	His	Val	Glu 70	Glu	Pro	Ala	Phe	Leu	Pro 75	Asp	Pro	Asn	Asp	Gly 80
Ser	Leu	Tyr	Thr 85	Leu	Gly	Ser	Lys	Asn	Asn 90	Glu	Gly	Leu	Thr	Lys 95	Leu
Pro	Phe	Thr	Ile 100	Pro	Glu	Leu	Val	Gln	Ala 105	Ser	Pro	Cys	Arg 110	Ser	Ser
Asp	Gly	Ile 115	Leu	Tyr	Met	Gly	Lys 120	Lys	Gln	Asp	Ile	Trp 125	Tyr	Val	Ile
Asp 130	Leu	Leu	Thr	Gly	Glu	Lys 135	Gln	Gln	Thr	Leu	Ser 140	Ser	Ala	Phe	Ala
Asp 145	Ser	Leu	Cys	Pro 150	Ser	Thr	Ser	Leu	Leu	Tyr 155	Leu	Gly	Arg	Thr	Glu 160
Tyr	Thr	Ile	Thr 165	Met	Tyr	Asp	Thr	Lys	Thr 170	Arg	Glu	Leu	Arg	Trp 175	Asn
Ala	Thr	Tyr	Phe 180	Asp	Tyr	Ala	Ala	Ser 185	Leu	Pro	Glu	Asp	Glu 190	Gly	Asp
Tyr	Lys 195	Met	Ser	His	Phe	Val	Ser 200	Asn	Gly	Asp	Gly	Leu 205	Val	Val	Thr
Val 210	Asp	Ser	Glu	Ser	Gly	Asp 215	Val	Leu	Trp	Ile	Gln 220	Asn	Tyr	Ala	Ser
Pro 225	Val	Val	Ala	Phe 230	Tyr	Val	Trp	Gln	Arg	Glu 235	Gly	Leu	Arg	Lys	Val 240
Met	His	Ile	Asn 245	Val	Ala	Val	Glu	Thr	Leu 250	Arg	Tyr	Leu	Thr	Phe 255	Met
Ser	Gly	Glu	Val 260	Gly	Arg	Ile	Thr	Lys 265	Trp	Lys	Tyr	Pro	Phe 270	Pro	Lys
Glu	Thr 275	Glu	Ala	Lys	Ser	Lys	Leu 280	Thr	Pro	Thr	Leu	Tyr 285	Val	Gly	Lys
Tyr 290	Ser	Thr	Ser	Leu	Tyr	Ala 295	Ser	Pro	Ser	Met	Val 300	His	Glu	Gly	Val

Ala Val Val Pro Arg Gly Ser Thr Leu Pro Leu Leu Glu Gly Pro Gln
305 310 315 320

Thr Asp Gly Val Thr Ile Gly Asp Lys Gly Glu Cys Val Ile Thr Pro
325 330 335

Ser Thr Asp Val Lys Phe Asp Pro Gly Leu Lys Ser Lys Asn Lys Leu
340 345 350

Asn Tyr Leu Arg Asn Tyr Trp Leu Leu Ile Gly His His Glu Thr Pro
355 360 365

Leu Ser Ala Ser Thr Lys Met Leu Glu Arg Phe Pro Asn Asn Leu Pro
370 375 380

Lys His Arg Glu Asn Val Ile Pro Ala Asp Ser Glu Lys Lys Ser Phe
385 390 395 400

Glu Glu Thr Leu Leu Gln Met Thr Ser Val Phe Ser Trp Ile Leu Asn
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Leu Pro Ser Lys Glu Glu Val Phe Ala Phe Leu Arg Ile Phe Glu Lys
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<210> 93
<211> 232
<212> PRT
<213> Homo sapiens
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<400> 93

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35 40 45

Ser Pro Glu Thr Ser Arg Ser Leu Pro Ala Pro Gln Asp Asn Asp Phe
50 55 60

Leu Ser Arg Lys Ala Gln Asp Cys Tyr Phe Met Lys Leu His His Cys
65 70 75 80

Pro Gly Asn His Ser Trp Asp Ser Thr Ile Ser Gly Ser Gln Arg Ala
85 90 95

Ala Phe Cys Asp His Lys Thr Thr Pro Cys Ser Ser Ala Ile Ile Asn
100 105 110

Pro Leu Ser Thr Ala Gly Asn Ser Glu Arg Leu Gln Pro Gly Ile Ala

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145					150					155					160
Ile	Met	Lys	Glu	Asp	Tyr	Glu	Leu	Val	Ser	Thr	Lys	Pro	Thr	Arg	Thr
				165					170					175	
Ser	Lys	Val	Arg	Gln	Leu	Leu	Asp	Thr	Thr	Asp	Ile	Gln	Gly	Glu	Glu
			180					185					190		
Phe	Ala	Lys	Val	Ile	Val	Gln	Lys	Leu	Lys	Asp	Asn	Lys	Gln	Met	Gly
		195					200					205			
Leu	Gln	Pro	Tyr	Pro	Glu	Ile	Leu	Val	Val	Ser	Arg	Ser	Pro	Ser	Leu
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Met	Glu	Lys	Ser	Lys	Ala	Thr	Pro	Ala	Ala	Arg	Ala	Ser	Lys	Lys	Ile
			20					25					30		
Leu	Leu	Pro	Glu	Pro	Ser	Ile	Arg	Ser	Val	Met	Gln	Lys	Tyr	Leu	Glu
		35					40					45			
Asp	Arg	Gly	Glu	Val	Thr	Phe	Glu	Lys	Ile	Phe	Ser	Gln	Lys	Leu	Gly
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Tyr	Leu	Leu	Phe	Arg	Asp	Phe	Cys	Leu	Asn	His	Leu	Glu	Glu	Ala	Arg
65				70						75				80	
Pro	Leu	Val	Glu	Phe	Tyr	Glu	Glu	Ile	Lys	Lys	Tyr	Glu	Lys	Leu	Glu
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Thr	Glu	Glu	Glu	Arg	Val	Ala	Arg	Ser	Arg	Glu	Ile	Phe	Asp	Ser	Tyr
			100				105						110		
Ile	Met	Lys	Glu	Leu	Leu	Ala	Cys	Ser	His	Pro	Phe	Ser	Lys	Ser	Ala
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Thr	Glu	His	Val	Gln	Gly	His	Leu	Gly	Lys	Lys	Gln	Val	Pro	Pro	Asp
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 130 135 140
 145 150 155 160
 165 170 175
 180 185 190
 195 200 205
 210 215 220
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 35 40 45
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 65 70 75 80
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 100 105 110
 115 120 125
 130 135 140

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145 150 155 160

Val Phe Gln Lys Phe Ile Glu Arg Val Ala Leu Ala Ala Gly Ala Ala
165 170 175

Thr Leu Pro Ala Val Pro Ser Cys Pro Asn Pro Gln His Pro Gly Ser
180 185 190

Gly Thr Thr Ala Arg His Leu Gln Val Gly Pro Tyr Trp Pro Arg Leu
195 200 205

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<210> 95
<211> 454
<212> PRT
<213> Homo sapiens

<400> 95

Met Gly Leu Val Ser Ser Lys Lys Pro Asp Lys Glu Lys Pro Ile Lys
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Glu Lys Asp Lys Gly Gln Trp Ser Pro Leu Lys Val Ser Ala Gln Asp
20 25 30

Lys Asp Ala Pro Pro Leu Pro Pro Leu Val Val Phe Asn His Leu Thr
35 40 45

Pro Pro Pro Pro Asp Glu His Leu Asp Glu Asp Lys His Phe Val Val
50 55 60

Ala Leu Tyr Asp Tyr Thr Ala Met Asn Asp Arg Asp Leu Gln Met Leu
65 70 75 80

Lys Gly Glu Lys Leu Gln Val Leu Lys Gly Thr Gly Asp Trp Trp Leu
85 90 95

Ala Arg Ser Leu Val Thr Gly Arg Glu Gly Tyr Val Pro Ser Asn Phe
100 105 110

Val Ala Arg Val Glu Ser Leu Glu Met Glu Arg Trp Phe Phe Arg Ser
115 120 125

Gln Gly Arg Lys Glu Ala Glu Arg Gln Leu Leu Ala Pro Ile Asn Lys
130 135 140

Ala Gly Ser Phe Leu Ile Arg Glu Ser Glu Thr Asn Lys Gly Ala Phe
145 150 155 160

Ser Leu Ser Val Lys Asp Val Thr Thr Gln Gly Glu Leu Ile Lys His
165 170 175

Tyr Lys Ile Arg Cys Leu Asp Glu Gly Gly Tyr Tyr Ile Ser Pro Arg
180 185 190

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Ile Thr Phe Pro Ser Leu Gln Ala Leu Val Gln His Tyr Ser Ser Tyr
195 200 205

Tyr Lys Asn Asn Met Lys Val Ala Ile Lys Thr Leu Lys Glu Gly Thr
210 215 220

Met Ser Pro Glu Ala Phe Leu Gly Glu Ala Asn Val Met Lys Ala Leu
225 230 235 240

Gln His Glu Arg Leu Val Arg Leu Tyr Ala Val Val Thr Lys Glu Pro
245 250 255

Ile Tyr Ile Val Thr Glu Tyr Met Ala Arg Gly Cys Leu Leu Asp Phe
260 265 270

Leu Lys Thr Asp Glu Gly Ser Arg Leu Ser Leu Pro Arg Leu Ile Asp
275 280 285

Met Ser Ala Gln Ile Ala Glu Gly Met Ala Tyr Ile Glu Arg Met Asn
290 295 300

Ser Ile His Arg Asp Leu Arg Ala Ala Asn Ile Leu Val Ser Glu Ala
305 310 315 320

Leu Cys Cys Lys Ile Ala Asp Phe Gly Leu Ala Arg Ile Ile Asp Ser
325 330 335

Glu Tyr Thr Ala Gln Glu Gly Ala Lys Phe Pro Ile Lys Trp Thr Ala
340 345 350

Pro Glu Ala Ile His Phe Gly Val Phe Thr Ile Lys Ala Asp Val Trp
355 360 365

Ser Phe Gly Val Leu Leu Met Glu Val Val Thr Tyr Gly Arg Val Pro
370 375 380

Tyr Pro Gly Met Ser Asn Pro Glu Val Ile Arg Asn Leu Glu Arg Gly
385 390 395 400

Tyr Arg Met Pro Arg Pro Asp Thr Cys Pro Pro Glu Leu Tyr Arg Gly
405 410 415

Val Ile Ala Glu Cys Trp Arg Ser Arg Pro Glu Glu Arg Pro Thr Phe
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Glu Phe Leu Gln Ser Val Leu Glu Asp Phe Tyr Thr Ala Thr Glu Arg
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Gln Tyr Glu Leu Gln Pro
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<210> 96
<211> 82
<212> PRT
<213> Homo sapiens

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Lys Lys Ile Arg Leu Asp Thr Glu Thr Glu Gly Val Pro Ser Thr Ala
35 40 45
Ile Arg Glu Ile Ser Leu Leu Lys Glu Leu Asn His Pro Asn Ile Val
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Lys Leu Leu Asp Val Ile His Thr Glu Asn Lys Asn Ile Ser Leu Lys
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Glu Gly

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<211> 118

<212> PRT

<213> Homo sapiens

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35 40 45
Lys Val Ile Asp Val Thr Gly Gly Gly Ser Phe Ser Pro Glu Glu Val
50 55 60
Arg Glu Leu Arg Glu Ala Thr Leu Lys Glu Val Asp Ile Leu Arg Lys
65 70 75 80
Val Ser Gly His Pro Asn Ile Ser Ile Gln Leu Lys Asp Thr Tyr Glu
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Leu Phe Asp Leu Pro His
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<213> Homo sapiens

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 35 40 45
 Lys Asp Gly Ile Pro Ile Ser Ser Leu Arg Glu Ile Thr Leu Leu Leu
 50 55 60
 Arg Leu Arg His Pro Asn Ile Val Glu Leu Lys Glu Val Val Val Gly
 65 70 75 80
 Asn His Leu Glu Ser Ile Phe Leu Val Met Gly Tyr Cys Glu Gln Asp
 85 90 95
 Leu Ala Ser Leu Leu Glu Asn Met Pro Thr Pro Phe Ser Glu Ala Gln
 100 105 110
 Val Lys Cys Ile Val Leu Gln Val Leu Arg Gly Leu Gln Tyr Leu His
 115 120 125
 Arg Asn Phe Ile Ile His Arg Asp Leu Lys Val Ser Asn Leu Leu Met
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 Trp Ser Leu Asp Gly Thr Trp
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<210> 99
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 <213> Homo sapiens

<400> 99

Met Ser Ser Ala Gly Gly Val Ser Arg Arg Leu Ala Ala Val Arg Ser
 1 5 10 15
 Thr Val Leu Cys Arg Ala Val Gly Cys Ile Leu Ala Glu Leu Leu Ala
 20 25 30
 His Arg Pro Leu Leu Pro Gly Thr Ser Glu Ile His Gln Ile Asp Leu
 35 40 45
 Ile Val Gln Leu Leu Gly Thr Pro Ser Glu Asn Ile Trp Pro Gly Phe
 50 55 60
 Ser Lys Leu Pro Leu Val Gly Gln Tyr Ser Leu Arg Lys Gln Pro Tyr
 65 70 75 80
 Asn Asn Leu Lys His Lys Phe Pro Trp Leu Ser Glu Ala Gly Leu Arg

85

90

95

Leu Leu His Phe Leu Phe Met Tyr Asp Pro Lys Lys Arg Ala Thr Ala
 100 105 110

Gly Asp Cys Leu Glu Ser Ser Tyr Phe Lys Glu Lys Pro Leu Arg Leu
 115 120 125

Pro Ile Ser Gly Val Cys Glu Gly Cys Arg Glu Pro Gly
 130 135 140

<210> 100

<211> 119

<212> PRT

<213> Homo sapiens

<400> 100

Val Phe Leu Gly Arg Cys Arg Ser Val Lys Glu Phe Glu Lys Leu Asn
 1 5 10 15

Arg Ile Gly Glu Gly Thr Tyr Gly Ile Val Tyr Arg Ala Arg Asp Thr
 20 25 30

Gln Thr Asp Glu Ile Val Ala Leu Lys Lys Val Arg Met Asp Lys Glu
 35 40 45

Lys Asp Gly Ile Pro Ile Ser Ser Leu Arg Glu Ile Thr Leu Leu Leu
 50 55 60

Arg Leu Arg His Pro Asn Ile Leu Pro Ala Arg Ala Pro Trp Lys Gly
 65 70 75 80

Arg Ser Gly Gly Ser Ile Arg Gly Cys Arg Gly Leu Met Trp Ser Ser
 85 90 95

Ser Leu Cys Trp Lys Cys Ala Thr Thr Ala Ser Trp Glu Glu Trp Trp
 100 105 110

Val Gln Ser Pro Arg Cys Leu
 115

<210> 101

<211> 756

<212> PRT

<213> Homo sapiens

<400> 101

Met Gly Glu Ala Glu Lys Phe His Tyr Ile Tyr Ser Cys Asp Leu Asp
 1 5 10 15

Ile Asn Val Gln Leu Lys Ile Gly Ser Leu Glu Gly Lys Arg Glu Gln
 20 25 30

Lys Ser Tyr Lys Ala Val Leu Glu Asp Pro Met Leu Lys Phe Ser Gly
 35 40 45

Leu Tyr Gln Glu Thr Cys Ser Asp Leu Tyr Val Thr Cys Gln Val Phe
 50 55 60
 Ala Glu Gly Lys Pro Leu Ala Leu Pro Val Arg Thr Ser Tyr Lys Ala
 65 70 75 80
 Phe Ser Thr Arg Trp Asn Trp Asn Glu Trp Leu Lys Leu Pro Val Lys
 85 90 95
 Tyr Pro Asp Leu Pro Arg Asn Ala Gln Val Ala Leu Thr Ile Trp Asp
 100 105 110
 Val Tyr Gly Pro Gly Lys Ala Val Pro Val Gly Gly Thr Thr Val Ser
 115 120 125
 Leu Phe Gly Lys Tyr Gly Met Phe Arg Gln Gly Met His Asp Leu Lys
 130 135 140
 Val Trp Pro Asn Val Glu Ala Asp Gly Ser Glu Pro Thr Lys Thr Pro
 145 150 155 160
 Gly Arg Thr Ser Ser Thr Leu Ser Glu Asp Gln Met Ser Arg Leu Ala
 165 170 175
 Lys Leu Thr Lys Ala His Arg Gln Gly His Met Val Lys Val Asp Trp
 180 185 190
 Leu Asp Arg Leu Thr Phe Arg Glu Ile Glu Met Ile Asn Glu Ser Val
 195 200 205
 Lys Arg Ser Ser Asn Phe Met Tyr Leu Met Gly Gly Phe Arg Cys Val
 210 215 220
 Lys Cys Asp Asp Lys Glu Tyr Gly Ile Val Tyr Tyr Glu Lys Asp Gly
 225 230 235 240
 Asp Glu Ser Ser Pro Ile Leu Thr Ser Phe Glu Leu Val Lys Val Pro
 245 250 255
 Asp Pro Gln Met Ser Leu Glu Asn Leu Val Glu Ser Lys His His Asn
 260 265 270
 Leu Pro Arg Ser Leu Arg Ser Gly Pro Ser Asp His Asp Leu Lys Pro
 275 280 285
 Tyr Pro Ser Pro Arg Asp Gln Leu Lys Asn Ile Val Ser Tyr Pro Pro
 290 295 300
 Ser Lys Pro Pro Thr Tyr Glu Glu Gln Asp Leu Val Trp Glu Phe Arg
 305 310 315 320
 Tyr Tyr Leu Thr Asn Gln Asp Lys Ala Leu Thr Lys Ile Leu Thr Ser
 325 330 335
 Val Ile Trp Asp Leu Pro Gln Glu Ala Lys Gln Ala Leu Ala Leu Leu
 340 345 350

Gly Lys Trp Asn Pro Met Asp Val Glu Asp Ser Leu Glu Leu Ile Ser
355 360 365

Ser His Tyr Thr Asn Pro Thr Val Arg Arg Tyr Ala Val Ala Arg Leu
370 375 380

Arg Gln Ala Asp Asp Glu Asp Leu Leu Met Tyr Leu Leu Gln Leu Val
385 390 395 400

Gln Ala Leu Lys Tyr Glu Asn Phe Asp Asp Ile Lys Asn Gly Leu Glu
405 410 415

Pro Thr Lys Lys Asp Ser Gln Ser Ser Val Ser Glu Asn Val Ser Asn
420 425 430

Ser Gly Ile Asn Ser Ala Glu Ile Asp Ser Ser Gln Ile Ile Thr Ser
435 440 445

Pro Leu Pro Ser Val Ser Ser Pro Pro Pro Ala Ser Lys Thr Lys Glu
450 455 460

Val Pro Asp Gly Glu Asn Leu Glu Gln Asp Leu Cys Thr Phe Leu Ile
465 470 475 480

Ser Arg Ala Cys Lys Asn Ser Thr Leu Ala Asn Tyr Leu Tyr Trp Tyr
485 490 495

Val Ile Val Glu Cys Glu Asp Gln Asp Thr Gln Gln Arg Asp Pro Lys
500 505 510

Thr His Glu Met Tyr Leu Asn Val Met Arg Arg Phe Ser Gln Ala Leu
515 520 525

Leu Lys Gly Asp Lys Ser Val Arg Val Met Arg Ser Leu Leu Ala Ala
530 535 540

Gln Gln Thr Phe Val Asp Arg Leu Val His Leu Met Lys Ala Val Gln
545 550 555 560

Arg Glu Ser Gly Asn Arg Lys Lys Lys Asn Glu Arg Leu Gln Ala Leu
565 570 575

Leu Gly Asp Asn Glu Lys Met Asn Leu Ser Asp Val Glu Leu Ile Pro
580 585 590

Leu Pro Leu Glu Pro Gln Val Lys Ile Arg Gly Ile Ile Pro Glu Thr
595 600 605

Ala Thr Leu Phe Lys Ser Ala Leu Met Pro Ala Gln Leu Phe Phe Lys
610 615 620

Thr Glu Asp Gly Gly Lys Tyr Pro Val Ile Phe Lys His Gly Asp Asp
625 630 635 640

Leu Arg Gln Asp Gln Leu Ile Leu Gln Ile Ile Ser Leu Met Asp Lys
645 650 655

Leu Leu Arg Lys Glu Asn Leu Asp Leu Lys Leu Thr Pro Tyr Lys Val
660 665 670

Leu Ala Thr Ser Thr Lys His Gly Phe Met Gln Phe Ile Gln Ser Val
675 680 685

Pro Val Ala Glu Val Leu Asp Thr Glu Gly Ser Ile Gln Asn Phe Phe
690 695 700

Arg Lys Tyr Ala Pro Ser Glu Asn Gly Pro Asn Gly Ile Ser Ala Glu
705 710 715 720

Val Met Asp Thr Tyr Val Lys Ser Cys Ala Gly Tyr Cys Val Ile Thr
725 730 735

Tyr Ile Leu Gly Val Gly Asp Arg His Leu Asp Asn Leu Leu Leu Thr
740 745 750

Lys Thr Gly Gly
755

<210> 102
<211> 508
<212> PRT
<213> Homo sapiens

<400> 102

Met Gly Glu Ala Glu Lys Phe His Tyr Ile Tyr Ser Cys Asp Leu Asp
1 5 10 15

Ile Asn Val Gln Leu Lys Ile Gly Ser Leu Glu Gly Lys Arg Glu Gln
20 25 30

Lys Ser Tyr Lys Ala Val Leu Glu Asp Pro Met Leu Lys Phe Ser Gly
35 40 45

Leu Tyr Gln Glu Thr Cys Ser Asp Leu Tyr Val Thr Cys Gln Val Phe
50 55 60

Ala Glu Gly Lys Pro Leu Ala Leu Pro Val Arg Thr Ser Tyr Lys Ala
65 70 75 80

Phe Ser Thr Arg Trp Asn Trp Asn Glu Trp Leu Lys Leu Pro Val Lys
85 90 95

Tyr Pro Asp Leu Pro Arg Asn Ala Gln Val Ala Leu Thr Ile Trp Asp
100 105 110

Val Tyr Gly Pro Gly Lys Ala Val Pro Val Gly Gly Thr Thr Val Ser
115 120 125

Leu Phe Gly Lys Tyr Gly Met Phe Arg Gln Gly Met His Asp Leu Lys
130 135 140

Val Trp Pro Asn Val Glu Ala Asp Gly Ser Glu Pro Thr Lys Thr Pro

145		150		155		160
Gly Arg Thr Ser Ser Thr Leu Ser Glu Asp Gln Met Ser Arg Leu Ala						
	165			170		175
Lys Leu Thr Lys Ala His Arg Gln Gly His Met Val Lys Val Asp Trp						
	180			185		190
Leu Asp Arg Leu Thr Phe Arg Glu Ile Glu Met Ile Asn Glu Ser Val						
	195			200		205
Lys Arg Ser Ser Asn Phe Met Tyr Leu Met Gly Gly Phe Arg Cys Val						
	210			215		220
Lys Cys Asp Asp Lys Glu Tyr Gly Ile Val Tyr Tyr Glu Lys Asp Gly						
	225			230		235
Asp Glu Ser Ser Pro Ile Leu Thr Ser Phe Glu Leu Val Lys Val Pro						
	245			250		255
Asp Pro Gln Met Ser Leu Glu Asn Leu Val Glu Ser Lys His His Asn						
	260			265		270
Leu Pro Arg Ser Leu Arg Ser Gly Pro Ser Asp His Asp Leu Lys Pro						
	275			280		285
Tyr Pro Ser Pro Arg Asp Gln Leu Lys Asn Ile Val Ser Tyr Pro Pro						
	290			295		300
Ser Lys Pro Pro Thr Tyr Glu Glu Gln Asp Leu Val Trp Glu Phe Arg						
	305			310		315
Tyr Tyr Leu Thr Asn Gln Asp Lys Ala Leu Thr Lys Ile Leu Thr Ser						
	325			330		335
Val Ile Trp Asp Leu Pro Gln Glu Ala Lys Gln Ala Leu Ala Leu Leu						
	340			345		350
Gly Lys Trp Asn Pro Met Asp Val Glu Asp Ser Leu Glu Leu Ile Ser						
	355			360		365
Ser His Tyr Thr Asn Pro Thr Val Arg Arg Tyr Ala Val Ala Arg Leu						
	370			375		380
Arg Gln Ala Asp Asp Glu Asp Leu Leu Met Tyr Leu Leu Gln Leu Val						
	385			390		395
Gln Ala Leu Lys Tyr Glu Asn Phe Asp Asp Ile Lys Asn Gly Leu Glu						
	405			410		415
Pro Thr Lys Lys Asp Ser Gln Ser Ser Val Ser Glu Asn Val Ser Asn						
	420			425		430
Ser Gly Ile Asn Ser Ala Glu Ile Asp Ser Ser Gln Ile Ile Thr Ser						
	435			440		445
Pro Leu Pro Ser Val Ser Ser Pro Pro Pro Ala Ser Lys Thr Lys Glu						

450 455 460

Val Pro Asp Gly Glu Asn Leu Glu Gln Asp Leu Cys Thr Phe Leu Ile
 465 470 475 480

Ser Arg Ala Cys Lys Asn Ser Thr Leu Ala Asn Tyr Leu Tyr Trp Tyr
 485 490 495

Val Lys Ile Ile Phe Cys Leu Phe Ser Tyr Tyr Pro
 500 505

<210> 103
 <211> 140
 <212> PRT
 <213> Homo sapiens

<400> 103

Met Gly Asn Ala Ala Ala Ala Lys Lys Gly Ser Glu Gln Glu Ser Val
 1 5 10 15

Lys Glu Phe Leu Ala Lys Ala Lys Glu Asp Phe Leu Lys Lys Trp Glu
 20 25 30

Ser Pro Ala Gln Asn Thr Ala His Leu Asp Gln Phe Glu Arg Ile Lys
 35 40 45

Thr Leu Gly Thr Gly Ser Phe Gly Arg Val Met Leu Val Lys His Lys
 50 55 60

Glu Thr Gly Asn His Tyr Ala Met Lys Ile Leu Asp Lys Gln Lys Val
 65 70 75 80

Val Lys Leu Lys Gln Ile Glu His Thr Leu Asn Glu Lys Arg Ile Leu
 85 90 95

Gln Ala Val Asn Phe Pro Phe Leu Val Lys Leu Glu Phe Ser Phe Lys
 100 105 110

Asp Asn Ser Asn Leu Tyr Met Val Met Glu Tyr Val Pro Gly Gly Glu
 115 120 125

Met Phe Ser His Leu Arg Arg Ile Gly Arg Phe Arg
 130 135 140

<210> 104
 <211> 156
 <212> PRT
 <213> Homo sapiens

<400> 104

Met Val Val Phe Asn Gly Leu Leu Lys Ile Lys Ile Cys Glu Ala Val
 1 5 10 15

Ser Leu Lys Pro Thr Ala Trp Ser Leu Arg His Ala Val Gly Pro Arg
 20 25 30

Pro Gln Thr Phe Leu Leu Asp Pro Tyr Ile Ala Leu Asn Val Asp Asp
35 40 45

Ser Arg Ile Gly Gln Thr Ala Thr Lys Gln Lys Thr Asn Ser Pro Ala
50 55 60

Trp His Asp Glu Phe Val Thr Asp Val Cys Asn Gly Arg Lys Ile Glu
65 70 75 80

Leu Ala Val Phe His Asp Ala Pro Ile Gly Tyr Asp Asp Phe Val Ala
85 90 95

Asn Cys Thr Ile Gln Phe Glu Glu Leu Leu Gln Asn Gly Ser Arg His
100 105 110

Phe Glu Asp Trp Ile Asp Leu Glu Pro Glu Gly Arg Val Tyr Val Ile
115 120 125

Ile Asp Leu Ser Gly Ser Ser Gly Glu Val Lys Ile Pro Asn Ser Ala
130 135 140

Phe Cys Glu Arg Glu Arg Val Glu Met Arg His Ser
145 150 155

<210> 105

<211> 520

<212> PRT

<213> Homo sapiens

<400> 105

Met Ile Leu Ile Pro Arg Met Leu Leu Val Leu Phe Leu Leu Leu Pro
1 5 10 15

Ile Leu Ser Ser Ala Lys Ala Gln Val Asn Pro Ala Ile Cys Arg Tyr
20 25 30

Pro Leu Gly Met Ser Gly Gly Gln Ile Pro Asp Glu Asp Ile Thr Ala
35 40 45

Ser Ser Gln Trp Ser Glu Ser Thr Ala Ala Lys Tyr Gly Arg Leu Asp
50 55 60

Ser Glu Glu Gly Asp Gly Ala Trp Cys Pro Glu Ile Pro Val Glu Pro
65 70 75 80

Asp Asp Leu Lys Glu Phe Leu Gln Ile Asp Leu His Thr Leu His Phe
85 90 95

Ile Thr Leu Val Gly Thr Gln Gly Arg His Ala Gly Gly His Gly Ile
100 105 110

Glu Phe Ala Pro Met Tyr Lys Ile Asn Tyr Ser Arg Asp Gly Thr Arg
115 120 125

Trp Ile Ser Trp Arg Asn Arg His Gly Lys Gln Val Leu Asp Gly Asn

435	440	445
Ser Asp Ser Ser Met Phe Asn Asn Asn Arg Ser Ser Ser Pro Ser Glu		
450	455	460
Gln Gly Ser Asn Ser Thr Tyr Asp Arg Ile Phe Pro Leu Arg Pro Asp		
465	470	475 480
Tyr Gln Glu Pro Ser Arg Leu Ile Arg Lys Leu Pro Glu Phe Ala Pro		
	485	490 495
Gly Glu Glu Glu Ser Gly Glu Asp Asp Val Val Glu Gln Gly Val Lys		
	500	505 510
Gly Glu Thr Ser Ala Ser Ile Arg		
	515	520
<210> 106		
<211> 284		
<212> PRT		
<213> Homo sapiens		
<400> 106		
Met Ala Asn Phe Gln Glu His Leu Ser Cys Ser Ser Ser Pro His Leu		
1	5	10 15
Pro Phe Ser Glu Ser Lys Thr Phe Asn Gly Leu Gln Asp Glu Leu Thr		
	20	25 30
Ala Met Gly Asn His Pro Ser Pro Lys Leu Leu Glu Asp Gln Gln Glu		
	35	40 45
Lys Gly Met Val Arg Thr Glu Leu Ile Glu Ser Val His Ser Pro Val		
	50	55 60
Thr Thr Thr Val Leu Thr Ser Val Ser Glu Asp Ser Arg Asp Gln Phe		
65	70	75 80
Glu Asn Ser Val Leu Gln Leu Arg Glu His Asp Glu Ser Glu Thr Ala		
	85	90 95
Val Ser Gln Gly Asn Ser Asn Thr Val Asp Gly Glu Ser Thr Ser Gly		
	100	105 110
Thr Glu Asp Ile Lys Ile Gln Phe Ser Arg Ser Gly Ser Gly Ser Gly		
	115	120 125
Gly Phe Leu Glu Gly Leu Phe Gly Cys Leu Arg Pro Val Trp Asn Ile		
	130	135 140
Ile Gly Lys Ala Tyr Ser Thr Asp Tyr Lys Phe Met Gln Gln Asp Thr		
145	150	155 160
Trp Glu Val Pro Phe Glu Glu Ile Ser Glu Leu Gln Trp Leu Gly Ser		
	165	170 175

Gly Ala Gln Gly Ala Val Phe Leu Gly Lys Phe Arg Ala Glu Glu Val
 180 185 190
 Ala Ile Lys Lys Val Arg Glu Gln Asn Glu Thr Asp Ile Lys His Leu
 195 200 205
 Arg Lys Leu Lys His Pro Asn Ile Ile Ala Phe Lys Gly Val Cys Thr
 210 215 220
 Gln Ala Pro Cys Tyr Cys Ile Ile Met Glu Tyr Cys Ala His Gly Gln
 225 230 235 240
 Leu Tyr Glu Val Leu Arg Ala Gly Arg Lys Ile Thr Pro Arg Leu Leu
 245 250 255
 Val Asp Trp Ser Thr Gly Ile Ala Ser Gly Met Asn Tyr Leu His Leu
 260 265 270
 His Lys Ile Ile His Arg Asp Leu Lys Ser Pro Lys
 275 280

<210> 107
 <211> 185
 <212> PRT
 <213> Homo sapiens

<400> 107

Met Cys Gly Gln Arg Trp Ile His Asn Phe Thr Cys Leu Ala Phe Leu
 1 5 10 15
 Phe His Thr Leu Lys Ser Gly Asn Lys Ser Val His Leu Arg Lys Ala
 20 25 30
 Ser Ser Pro Asn Leu His Arg Arg Gln Trp Glu Lys Asn Val Pro Asn
 35 40 45
 Thr Ala Leu Thr Ala Leu Glu Asn Ala Ser Ile Leu Thr Ser Ser Leu
 50 55 60
 Thr Ala Glu Asp Asp Arg Gly Gly Ser Val Ile Lys Tyr Ser Lys Asn
 65 70 75 80
 Thr Thr Arg Lys Gln Trp Leu Lys Glu Thr Pro Asp Thr Leu Leu Asn
 85 90 95
 Ile Leu Lys Asn Ala Asp Leu Ser Leu Ala Phe Gln Thr Tyr Thr Ile
 100 105 110
 Tyr Arg Pro Gly Ser Glu Gly Phe Leu Lys Gly Pro Leu Ser Glu Glu
 115 120 125
 Thr Glu Ala Ser Asp Ser Val Asp Gly Gly His Asp Ser Val Ile Leu
 130 135 140
 Asp Pro Glu Arg Leu Glu Pro Gly Leu Asp Glu Glu Asp Thr Asp Phe
 145 150 155 160

Glu Glu Glu Asp Asp Asn Pro Asp Trp Val Ser Glu Leu Lys Lys Arg
165 170 175

Ala Gly Trp Gln Gly Leu Cys Asp Arg
180 185

<210> 108
<211> 83
<212> PRT
<213> Homo sapiens

<400> 108

Met Ala Pro Pro Ser Glu Glu Thr Pro Leu Ile Pro Gln Arg Ser Cys
1 5 10 15

Ser Leu Leu Ser Thr Glu Ala Gly Ala Leu His Val Leu Leu Pro Ala
20 25 30

Arg Gly Pro Gly Pro Pro Gln Arg Leu Ser Phe Ser Phe Gly Val Pro
35 40 45

Val Arg Pro Val Gly Ala Asn Gly Pro Pro Leu Thr Ser Gly Phe Leu
50 55 60

Gly Gly Trp Ala Glu Ala Ser Val Gln Arg Gly Leu Trp Lys Cys Leu
65 70 75 80

Leu Thr Glu

<210> 109
<211> 213
<212> PRT
<213> Homo sapiens

<400> 109

Met Ala Glu Ser Ala Gly Ala Ser Ser Phe Phe Pro Leu Val Val Leu
1 5 10 15

Leu Leu Ala Gly Ser Gly Gly Ser Gly Pro Arg Gly Val Gln Ala Leu
20 25 30

Leu Cys Ala Cys Thr Ser Cys Leu Gln Ala Asn Tyr Thr Cys Glu Thr
35 40 45

Asp Gly Ala Cys Met Val Ser Ile Phe Asn Leu Asp Gly Met Glu His
50 55 60

His Val Arg Thr Cys Ile Pro Lys Val Glu Leu Val Pro Ala Gly Lys
65 70 75 80

Pro Phe Tyr Cys Leu Ser Ser Glu Asp Leu Arg Asn Thr His Cys Cys
85 90 95

Tyr Thr Asp Tyr Cys Asn Arg Ile Asp Leu Arg Val Pro Ser Gly His
100 105 110

Leu Lys Glu Pro Glu His Pro Ser Met Trp Gly Pro Val Glu Leu Val
115 120 125

Gly Ile Ile Ala Gly Pro Val Phe Leu Leu Phe Leu Ile Ile Ile Ile
130 135 140

Val Phe Leu Val Ile Asn Tyr His Gln Arg Val Tyr His Asn Arg Gln
145 150 155 160

Arg Leu Asp Met Glu Asp Pro Ser Cys Glu Met Cys Leu Ser Lys Asp
165 170 175

Lys Thr Leu Gln Asp Leu Val Tyr Asp Leu Ser Thr Ser Gly Ser Gly
180 185 190

Ser Gly Thr Lys Phe Phe Arg Ala Ser Cys Leu Trp Leu Ala Phe Ile
195 200 205

Ser Phe Pro Ala Gly
210

<210> 110
<211> 383
<212> PRT
<213> Homo sapiens

<400> 110

Met Asp Glu Gln Glu Ala Leu Asn Ser Ile Met Asn Asp Leu Val Ala
1 5 10 15

Leu Gln Met Asn Arg Arg His Arg Met Pro Gly Tyr Glu Thr Met Lys
20 25 30

Asn Lys Asp Thr Gly His Ser Asn Arg Gln Ser Asp Val Arg Ile Lys
35 40 45

Phe Glu His Asn Gly Glu Arg Arg Ile Ile Ala Phe Ser Arg Pro Val
50 55 60

Lys Tyr Glu Asp Val Glu His Lys Val Thr Thr Val Phe Gly Gln Pro
65 70 75 80

Leu Asp Leu His Tyr Met Asn Asn Glu Leu Ser Ile Leu Leu Lys Asn
85 90 95

Gln Asp Asp Leu Asp Lys Ala Ile Asp Ile Leu Asp Arg Ser Ser Ser
100 105 110

Met Lys Ser Leu Arg Ile Leu Leu Leu Ser Gln Asp Arg Asn His Asn
115 120 125

Ser Ser Ser Pro His Ser Glu Val Ser Arg Gln Val Arg Ile Lys Ala
130 135 140

Ser Gln Ser Ala Gly Asp Ile Asn Thr Ile Tyr Gln Pro Pro Glu Pro
 145 150 155 160
 Arg Ser Arg His Leu Ser Val Ser Ser Gln Asn Pro Gly Arg Ser Ser
 165 170 175
 Pro Pro Pro Gly Tyr Val Pro Glu Arg Gln Gln His Ile Ala Arg Gln
 180 185 190
 Gly Ser Tyr Thr Ser Ile Asn Ser Glu Gly Glu Phe Ile Pro Glu Thr
 195 200 205
 Ser Glu Gln Cys Met Leu Asp Pro Leu Ser Ser Ala Glu Asn Ser Leu
 210 215 220
 Ser Gly Ser Cys Gln Ser Leu Asp Arg Ser Ala Asp Ser Pro Ser Phe
 225 230 235 240
 Arg Lys Ser Arg Met Ser Arg Ala Gln Ser Phe Pro Asp Asn Arg Gln
 245 250 255
 Glu Tyr Ser Asp Arg Glu Thr Gln Leu Tyr Asp Lys Gly Val Lys Gly
 260 265 270
 Gly Thr Tyr Pro Arg Arg Tyr His Val Ser Val His His Lys Asp Tyr
 275 280 285
 Ser Asp Gly Arg Arg Thr Phe Pro Arg Ile Arg Arg His Gln Gly Asn
 290 295 300
 Leu Phe Thr Leu Val Pro Ser Ser Arg Ser Leu Ser Thr Asn Gly Glu
 305 310 315 320
 Asn Met Gly Leu Ala Val Gln Tyr Leu Asp Pro Arg Gly Arg Leu Arg
 325 330 335
 Ser Ala Asp Ser Glu Asn Ala Leu Ser Val Gln Glu Arg Asn Val Pro
 340 345 350
 Thr Lys Cys Glu Glu Leu Ser Leu Ala Arg Arg Arg Leu Pro Arg Trp
 355 360 365
 Ser Gln Thr Ser Tyr Gly Gly Lys Gln Leu Gly Pro Trp Asp Pro
 370 375 380

<210> 111

<211> 414

<212> PRT

<213> Homo sapiens

<400> 111

Met Asp Glu Gln Glu Ala Leu Asn Ser Ile Met Asn Asp Leu Val Ala
 1 5 10 15

Leu Gln Met Asn Arg Arg His Arg Met Pro Gly Tyr Glu Thr Met Lys

20							25					30						
Asn	Lys	Asp	Thr	Gly	His	Ser	Asn	Arg	Gln	Lys	Lys	His	Asn	Ser	Ser			
35							40					45						
Ser	Ser	Ala	Leu	Leu	Asn	Ser	Pro	Thr	Val	Thr	Thr	Ser	Ser	Cys	Ala			
50							55					60						
Gly	Ala	Ser	Glu	Lys	Lys	Lys	Phe	Leu	Ser	Asp	Val	Arg	Ile	Lys	Phe			
65							70					75					80	
Glu	His	Asn	Gly	Glu	Arg	Arg	Ile	Ile	Ala	Phe	Ser	Arg	Pro	Val	Lys			
85							90					95						
Tyr	Glu	Asp	Val	Glu	His	Lys	Val	Thr	Thr	Val	Phe	Gly	Gln	Pro	Leu			
100							105					110						
Asp	Leu	His	Tyr	Met	Asn	Asn	Glu	Leu	Ser	Ile	Leu	Leu	Lys	Asn	Gln			
115							120					125						
Asp	Asp	Leu	Asp	Lys	Ala	Ile	Asp	Ile	Leu	Asp	Arg	Ser	Ser	Ser	Met			
130							135					140						
Lys	Ser	Leu	Arg	Ile	Leu	Leu	Leu	Ser	Gln	Asp	Arg	Asn	His	Asn	Ser			
145							150					155					160	
Ser	Ser	Pro	His	Ser	Glu	Val	Ser	Arg	Gln	Val	Arg	Ile	Lys	Ala	Ser			
165							170					175						
Gln	Ser	Ala	Gly	Asp	Ile	Asn	Thr	Ile	Tyr	Gln	Pro	Pro	Glu	Pro	Arg			
180							185					190						
Ser	Arg	His	Leu	Ser	Val	Ser	Ser	Gln	Asn	Pro	Gly	Arg	Ser	Ser	Pro			
195							200					205						
Pro	Pro	Gly	Tyr	Val	Pro	Glu	Arg	Gln	Gln	His	Ile	Ala	Arg	Gln	Gly			
210							215					220						
Ser	Tyr	Thr	Ser	Ile	Asn	Ser	Glu	Gly	Glu	Phe	Ile	Pro	Glu	Thr	Ser			
225							230					235					240	
Glu	Gln	Cys	Met	Leu	Asp	Pro	Leu	Ser	Ser	Ala	Glu	Asn	Ser	Leu	Ser			
245							250					255						
Gly	Ser	Cys	Gln	Ser	Leu	Asp	Arg	Ser	Ala	Asp	Ser	Pro	Ser	Phe	Arg			
260							265					270						
Lys	Ser	Arg	Met	Ser	Arg	Ala	Gln	Ser	Phe	Pro	Asp	Asn	Arg	Gln	Glu			
275							280					285						
Tyr	Ser	Asp	Arg	Glu	Thr	Gln	Leu	Tyr	Asp	Lys	Gly	Val	Lys	Gly	Gly			
290							295					300						
Thr	Tyr	Pro	Arg	Arg	Tyr	His	Val	Ser	Val	His	His	Lys	Asp	Tyr	Ser			
305							310					315					320	
Asp	Gly	Arg	Arg	Thr	Phe	Pro	Arg	Ile	Arg	Arg	His	Gln	Gly	Asn	Leu			

Gln Ile Thr Gln Gln Ser Thr Asn Gln Ser Arg Asn Pro Ala Thr Thr
 180 185 190

Asn Gln Thr Glu Phe Glu Arg Val Phe
 195 200

<210> 113
 <211> 125
 <212> PRT
 <213> Homo sapiens
 <400> 113

Met Ala Thr Ser Arg Tyr Glu Pro Val Ala Glu Ile Gly Val Gly Ala
 1 5 10 15

Tyr Gly Thr Val Tyr Lys Ala Arg Asp Pro His Ser Gly His Phe Cys
 20 25 30

Ala Leu Lys Ser Val Arg Val Pro Asn Gly Gly Gly Gly Gly Gly Gly
 35 40 45

Leu Pro Ile Ser Thr Val Arg Glu Val Ala Leu Leu Arg Arg Leu Glu
 50 55 60

Ala Phe Glu His Pro Asn Val Val Arg Leu Met Asp Val Cys Ala Thr
 65 70 75 80

Ser Arg Thr Asp Arg Glu Ile Lys Val Thr Leu Val Phe Glu His Val
 85 90 95

Asp Gln Asp Leu Arg Thr Tyr Leu Asp Lys Ala Pro Pro Pro Gly Leu
 100 105 110

Pro Ala Glu Thr Ile Lys Val Ser Gly Val Gly Arg His
 115 120 125

<210> 114
 <211> 45
 <212> PRT
 <213> Homo sapiens
 <400> 114

Met Ala Thr Ser Arg Tyr Glu Pro Val Ala Glu Ile Gly Val Gly Ala
 1 5 10 15

Tyr Gly Thr Val Tyr Lys Ala Arg Asp Pro His Ser Gly His Phe Cys
 20 25 30

Ala Leu Lys Ser Val Arg Val Pro Thr His Leu Ser Phe
 35 40 45

<210> 115
 <211> 160
 <212> PRT
 <213> Homo sapiens

<400> 115

```
Met Gly Val Cys Pro Gly Lys Thr Pro Phe Cys Ser Pro Lys Pro Gln
1          5          10          15

Gly Leu Ala Arg Gly His Trp Ser Arg Arg Arg Asp Ile Cys Val Thr
          20          25          30

Gly Pro Leu Pro Leu Glu Pro Arg Ala Val Tyr Cys Lys Asp Val Leu
          35          40          45

Asp Ile Glu Gln Phe Ser Thr Val Lys Gly Val Asn Leu Asp His Thr
          50          55          60

Asp Asp Asp Phe Tyr Ser Lys Phe Ser Thr Gly Ser Val Ser Ile Pro
65          70          75          80

Trp Gln Asn Glu Met Ile Glu Thr Glu Cys Phe Lys Glu Leu Asn Val
          85          90          95

Phe Gly Pro Asn Gly Thr Leu Pro Pro Asp Leu Asn Arg Asn His Pro
          100          105          110

Pro Glu Pro Pro Lys Lys Gly Leu Leu Gln Arg Leu Phe Lys Arg Gln
          115          120          125

His Gln Asn Asn Ser Lys Ser Ser Pro Ser Ser Lys Thr Ser Phe Asn
          130          135          140

His His Ile Asn Ser Asn His Val Ser Ser Asn Ser Thr Gly Ser Ser
145          150          155          160
```

<210> 116

<211> 300

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(300)

<223> "XAA" can be any amino acid

<400> 116

```
Met Pro Arg Ala Arg Met Pro Xaa Pro Arg Ala His Ser Lys Ala Gly
1          5          10          15

Cys Pro Cys Gly Cys Pro Arg Asp Pro Leu Thr Leu Leu Ser Pro Ser
          20          25          30

Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Val Lys Ile Pro Glu
          35          40          45

Gly Asp Leu Ile Arg Gly Arg Val Gly Thr Val Gly Tyr Met Ala Pro
          50          55          60
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Glu	Val	Leu	Asn	Asn	Gln	Arg	Tyr	Gly	Leu	Ser	Pro	Asp	Tyr	Trp	Gly	
65					70					75					80	
Leu	Gly	Cys	Leu	Ile	Tyr	Glu	Met	Ile	Glu	Gly	Gln	Ser	Pro	Phe	Arg	
				85					90					95		
Gly	Arg	Lys	Glu	Lys	Val	Lys	Arg	Glu	Glu	Val	Asp	Arg	Arg	Val	Leu	
			100					105					110			
Glu	Thr	Glu	Glu	Val	Tyr	Ser	His	Lys	Phe	Ser	Glu	Glu	Ala	Lys	Ser	
			115				120						125			
Ile	Cys	Lys	Met	Leu	Leu	Thr	Lys	Asp	Ala	Lys	Gln	Arg	Leu	Gly	Cys	
	130						135					140				
Gln	Glu	Glu	Gly	Ala	Ala	Glu	Val	Lys	Arg	His	Pro	Phe	Phe	Arg	Asn	
145					150					155					160	
Met	Asn	Phe	Lys	Arg	Leu	Glu	Ala	Gly	Met	Leu	Asp	Pro	Pro	Phe	Val	
				165					170					175		
Pro	Asp	Pro	Arg	Ala	Val	Tyr	Cys	Lys	Asp	Val	Leu	Asp	Ile	Glu	Gln	
			180					185					190			
Phe	Ser	Thr	Val	Lys	Gly	Val	Asn	Leu	Asp	His	Thr	Asp	Asp	Asp	Phe	
		195					200					205				
Tyr	Ser	Lys	Phe	Ser	Thr	Gly	Ser	Val	Ser	Ile	Pro	Trp	Gln	Asn	Glu	
	210					215					220					
Met	Ile	Glu	Thr	Glu	Cys	Phe	Lys	Glu	Leu	Asn	Val	Phe	Gly	Pro	Asn	
225					230					235					240	
Gly	Thr	Leu	Pro	Pro	Asp	Leu	Asn	Arg	Asn	His	Pro	Pro	Glu	Pro	Pro	
				245					250					255		
Lys	Lys	Gly	Leu	Leu	Gln	Arg	Leu	Phe	Lys	Arg	Gln	His	Gln	Asn	Asn	
			260					265					270			
Ser	Lys	Ser	Ser	Pro	Ser	Ser	Lys	Thr	Ser	Phe	Asn	His	His	Ile	Asn	
		275					280					285				
Ser	Asn	His	Val	Ser	Ser	Asn	Ser	Thr	Gly	Ser	Ser					
	290					295					300					

<210> 117

<211> 169

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(169)

<223> "XAA" can be any amino acid

<400> 117

Met Arg Met Pro Arg Ala Arg Met Pro Xaa Pro Arg Ala His Ser Lys
1 5 10 15
Ala Gly Cys Pro Cys Gly Cys Pro Arg Asp Pro Leu Thr Leu Leu Ser
20 25 30
Pro Ser Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Val Lys Ile
35 40 45
Pro Glu Gly Asp Leu Ile Arg Gly Arg Val Gly Thr Val Gly Tyr Met
50 55 60
Ala Pro Glu Val Leu Asn Asn Gln Arg Tyr Gly Leu Ser Pro Asp Tyr
65 70 75 80
Trp Gly Leu Gly Cys Leu Ile Tyr Glu Met Ile Glu Gly Gln Ser Pro
85 90 95
Phe Arg Gly Arg Lys Glu Lys Val Lys Arg Glu Glu Val Asp Arg Arg
100 105 110
Val Leu Glu Thr Glu Glu Val Tyr Ser His Lys Phe Ser Glu Glu Ala
115 120 125
Lys Ser Ile Cys Lys Met Val Ser Ser Trp Trp Pro Asp Ala Thr Leu
130 135 140
Lys Leu Val Ala Pro Ser Leu Gly Leu Ala Pro Val Cys Pro Gln Asn
145 150 155 160
Ser Lys Gln Ala Glu Gly Thr Gly Val
165

<210> 118

<211> 319

<212> PRT

<213> Homo sapiens

<400> 118

Met Ala Pro Phe Leu Arg Ile Ala Phe Asn Ser Tyr Glu Leu Gly Ser
1 5 10 15
Leu Gln Ala Glu Asp Glu Ala Asn Gln Pro Phe Cys Ala Val Lys Met
20 25 30
Lys Glu Ala Leu Ser Thr Glu Arg Gly Lys Thr Leu Val Gln Lys Lys
35 40 45
Pro Thr Met Tyr Pro Glu Trp Lys Ser Thr Phe Asp Ala His Ile Tyr
50 55 60
Glu Gly Arg Val Ile Gln Ile Val Leu Met Arg Ala Ala Glu Glu Pro
65 70 75 80

Val Ser Glu Val Thr Val Gly Val Ser Val Leu Ala Glu Arg Cys Lys
85 90 95

Lys Asn Asn Gly Lys Ala Glu Phe Trp Leu Asp Leu Gln Pro Gln Ala
100 105 110

Lys Val Leu Met Ser Val Gln Tyr Phe Leu Glu Asp Val Asp Cys Lys
115 120 125

Gln Ser Met Arg Ser Glu Asp Glu Ala Lys Phe Pro Thr Met Asn Arg
130 135 140

Arg Gly Ala Ile Lys Gln Ala Lys Ile His Tyr Ile Lys Asn His Glu
145 150 155 160

Phe Ile Ala Thr Phe Phe Gly Gln Pro Thr Phe Cys Ser Val Cys Lys
165 170 175

Asp Phe Val Trp Gly Leu Asn Lys Gln Gly Tyr Lys Cys Arg Gln Cys
180 185 190

Asn Ala Ala Ile His Lys Lys Cys Ile Asp Lys Ile Ile Gly Arg Cys
195 200 205

Thr Gly Thr Ala Ala Asn Ser Arg Asp Thr Ile Phe Gln Lys Glu Arg
210 215 220

Phe Asn Ile Asp Met Pro His Arg Phe Lys Val His Asn Tyr Met Ser
225 230 235 240

Pro Thr Phe Cys Asp His Cys Gly Ser Leu Leu Leu Pro Ala Pro His
245 250 255

Asp Lys His Gln Trp Asp Cys Gly Asp Phe Cys Cys Trp Pro Arg Pro
260 265 270

Cys Pro Gln Ser Val Leu Gly Cys Arg Leu Ala Gly Leu Ser Trp Tyr
275 280 285

Phe Leu Cys Glu Leu Cys Val Asn Leu Leu Phe Leu Cys Leu Arg Arg
290 295 300

Glu Ile Val Asn Pro Val Phe His Tyr Leu Asn Val Val Ile Tyr
305 310 315

<210> 119

<211> 236

<212> PRT

<213> Homo sapiens

<400> 119

Met Asp Glu Thr His Pro Gly Tyr Gly Lys Glu Val Asp Leu Glu Phe
1 5 10 15

Leu Val Ser Pro Ser Leu Pro Cys Leu Leu Ser Phe Ala Gly Ser Ala
20 25 30

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tcgccgttgt tctgtctcgt gccgccgccc tgggctgcat tgggttggtg gccacgccg 1080
gccaactcac cgcagtctgg cgccgcccag gagccgcccg cgctccctga accctagaac 1140
tgtcttcgac tccggggccc cgttggaaga ctgagtgccg ggggcacggc acagaagccg 1200
cgcccaccgc ctgccagttc acaaccgctc cgagcgtggg tctccgcca gctccagtcc 1260
tgtgatccgg gcccgccccc tagcggccgg ggagggaggg gccgggtccg cggccggcga 1320
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ggatgcgaag cggccgaatc agggttgggg gaggaaaagc cacggggcgg ggctttggcg 2040
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cagctgtgcc accgagcgtc gagaagaggg ggctgggctg gcagcgcgcg cggccatcct 2160
ccttccactg cgcctgcgca cgccacgcgc atccg 2195

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<210> 61
 <211> 1662
 <212> DNA
 <213> Homo sapiens

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<400> 61
ckwkkttact tcagacatgg gacggtctct gtagttacag tggggcatta agtaagggtg 60
tgtgtgttgc tggggatctg agaagtcgat ctttgagctg agcgtggtg aaggagaaac 120
aagccatgga aggaaagggt ccaagtggtc aggcgagagc ctccaggga aaggccttgg 180
gcagggtggga atcctgattt gttcctcagt caactacgcg aggcagaggc tcggaaccgg 240
gacctagagg cacacgtccg gcagttgcag gagcggatgg agttgctgca ggcagaggga 300

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Arg His Leu Val Pro Pro Asp Ser Asn Leu Phe Ser Lys Leu Trp Ala
 35 40 45
 Cys Gly Val Ile Leu Phe Thr Leu Leu Ala Gly Ser Pro Pro Phe Trp
 50 55 60
 His Arg Arg Gln Ile Leu Met Leu Arg Met Ile Met Glu Gly Gln Tyr
 65 70 75 80
 Gln Phe Ser Ser Pro Glu Trp Asp Asp Arg Ser Ser Thr Val Lys Asp
 85 90 95
 Leu Ile Ser Arg Leu Leu Gln Val Asp Pro Glu Ala Arg Leu Thr Ala
 100 105 110
 Glu Gln Ala Leu Gln His Pro Phe Phe Glu Arg Cys Glu Gly Ser Gln
 115 120 125
 Pro Trp Asn Leu Thr Pro Arg Gln Arg Phe Arg Val Ala Val Trp Thr
 130 135 140
 Val Leu Ala Ala Gly Arg Val Ala Leu Ser Thr His Arg Val Arg Pro
 145 150 155 160
 Leu Thr Lys Asn Ala Leu Leu Arg Asp Pro Tyr Ala Leu Arg Ser Val
 165 170 175
 Arg His Leu Ile Asp Asn Cys Ala Phe Arg Leu Tyr Gly His Trp Val
 180 185 190
 Lys Lys Gly Glu Gln Gln Asn Arg Ala Ala Leu Phe Gln His Arg Pro
 195 200 205
 Pro Gly Pro Phe Pro Ile Met Gly Pro Glu Glu Glu Gly Asp Ser Ala
 210 215 220
 Ala Ile Thr Glu Asp Glu Ala Val Leu Val Leu Gly
 225 230 235

<210> 120
 <211> 572
 <212> PRT
 <213> Homo sapiens

<400> 120

Met Ala Phe Cys Ala Lys Met Arg Ser Ser Lys Lys Thr Glu Val Asn
 1 5 10 15
 Leu Glu Ala Pro Glu Pro Gly Val Glu Val Ile Phe Tyr Leu Ser Asp
 20 25 30
 Arg Glu Pro Leu Arg Leu Gly Ser Gly Glu Tyr Thr Ala Glu Glu Leu
 35 40 45
 Cys Ile Arg Ala Ala Gln Ala Cys Arg Ile Ser Pro Leu Cys His Asn

50

55

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Leu Phe Ala Leu Tyr Asp Glu Asn Thr Lys Leu Trp Tyr Ala Pro Asn
65 70 75 80

Arg Thr Ile Thr Val Asp Asp Lys Met Ser Leu Arg Leu His Tyr Arg
85 90 95

Met Arg Phe Tyr Phe Thr Asn Trp His Gly Thr Asn Asp Asn Glu Gln
100 105 110

Ser Val Trp Arg His Ser Pro Lys Lys Gln Lys Asn Gly Tyr Glu Lys
115 120 125

Lys Lys Ile Pro Asp Ala Thr Pro Leu Leu Asp Ala Ser Ser Leu Glu
130 135 140

Tyr Leu Phe Ala Gln Gly Gln Tyr Asp Leu Val Lys Cys Leu Ala Pro
145 150 155 160

Ile Arg Asp Pro Lys Thr Glu Gln Asp Gly His Asp Ile Glu Asn Glu
165 170 175

Cys Leu Gly Met Ala Val Leu Ala Ile Ser His Tyr Ala Met Met Lys
180 185 190

Lys Met Gln Leu Pro Glu Leu Pro Lys Asp Ile Ser Tyr Lys Arg Tyr
195 200 205

Ile Pro Glu Thr Leu Asn Lys Ser Ile Arg Gln Arg Asn Leu Leu Thr
210 215 220

Arg Met Arg Ile Asn Asn Val Phe Lys Asp Phe Leu Lys Glu Phe Asn
225 230 235 240

Asn Lys Thr Ile Cys Asp Ser Ser Val Ser Thr His Asp Leu Lys Val
245 250 255

Lys Tyr Leu Ala Thr Leu Glu Thr Leu Thr Lys His Tyr Gly Ala Glu
260 265 270

Ile Phe Glu Thr Ser Met Leu Leu Ile Ser Ser Glu Asn Glu Met Asn
275 280 285

Trp Phe His Ser Asn Asp Gly Gly Asn Val Leu Tyr Tyr Glu Val Met
290 295 300

Val Thr Gly Asn Leu Gly Ile Gln Trp Arg His Lys Pro Asn Val Val
305 310 315 320

Ser Val Glu Lys Glu Lys Asn Lys Leu Lys Arg Lys Lys Leu Glu Asn
325 330 335

Lys Asp Lys Lys Asp Glu Glu Lys Asn Lys Ile Arg Glu Glu Trp Asn
340 345 350

Asn Phe Ser Phe Phe Pro Glu Ile Thr His Ile Val Ile Lys Glu Ser

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100

355					360					365						
Val	Val	Ser	Ile	Asn	Lys	Gln	Asp	Asn	Lys	Lys	Met	Glu	Leu	Lys	Leu	
370					375					380						
Ser	Ser	His	Glu	Glu	Ala	Leu	Ser	Phe	Val	Ser	Leu	Val	Asp	Gly	Tyr	
385					390					395					400	
Phe	Arg	Leu	Thr	Ala	Asp	Ala	His	His	Tyr	Leu	Cys	Thr	Asp	Val	Ala	
405					410					415						
Pro	Pro	Leu	Ile	Val	His	Asn	Ile	Gln	Asn	Gly	Cys	His	Gly	Pro	Ile	
420					425					430						
Cys	Thr	Glu	Tyr	Ala	Ile	Asn	Lys	Leu	Arg	Gln	Glu	Gly	Ser	Glu	Glu	
435					440					445						
Gly	Met	Tyr	Val	Leu	Arg	Trp	Ser	Cys	Thr	Asp	Phe	Asp	Asn	Ile	Leu	
450					455					460						
Met	Thr	Val	Thr	Cys	Phe	Glu	Lys	Ser	Glu	Gln	Val	Gln	Gly	Ala	Gln	
465					470					475					480	
Lys	Gln	Phe	Lys	Asn	Phe	Gln	Ile	Glu	Val	Gln	Lys	Gly	Arg	Tyr	Ser	
485					490					495						
Leu	His	Gly	Ser	Asp	Arg	Ser	Phe	Pro	Ser	Leu	Gly	Asp	Leu	Met	Ser	
500					505					510						
His	Leu	Lys	Lys	Gln	Ile	Leu	Arg	Thr	Asp	Asn	Ile	Ser	Phe	Met	Leu	
515					520					525						
Lys	Arg	Cys	Cys	Gln	Pro	Lys	Pro	Arg	Gly	Ser	Leu	Pro	Val	Pro	Glu	
530					535					540						
Pro	Gly	Cys	Ile	Pro	Ser	Val	Ile	Ala	Glu	Thr	Gln	Ile	Asp	Gln	Asn	
545					550					555					560	
Thr	Leu	Thr	Asp	Leu	Asn	Lys	Val	Asp	Pro	Pro	Pro					
565					570											

<210> 121
 <211> 311
 <212> PRT
 <213> Homo sapiens

<400> 121

Met	Gly	Cys	Val	Gln	Cys	Lys	Asp	Lys	Glu	Ala	Thr	Lys	Leu	Thr	Glu
1				5					10					15	
Glu	Arg	Asp	Gly	Ser	Leu	Asn	Gln	Ser	Ser	Gly	Tyr	Arg	Tyr	Gly	Thr
			20					25					30		
Asp	Pro	Thr	Pro	Gln	His	Tyr	Pro	Ser	Phe	Gly	Val	Thr	Ser	Ile	Pro
			35				40						45		

Asn Tyr Asn Asn Phe His Ala Ala Gly Gly Gln Gly Leu Thr Val Phe
 50 55 60
 Gly Gly Val Asn Ser Ser Ser His Thr Gly Thr Leu Arg Thr Arg Gly
 65 70 75 80
 Gly Thr Gly Val Thr Leu Phe Val Ala Leu Tyr Asp Tyr Glu Ala Arg
 85 90 95
 Thr Glu Asp Asp Leu Ser Phe His Lys Gly Glu Lys Phe Gln Ile Leu
 100 105 110
 Asn Ser Ser Glu Gly Asp Trp Trp Glu Ala Arg Ser Leu Thr Thr Gly
 115 120 125
 Glu Thr Gly Tyr Ile Pro Ser Asn Tyr Val Ala Pro Val Asp Ser Ile
 130 135 140
 Gln Ala Glu Glu Trp Tyr Phe Gly Lys Leu Gly Arg Lys Asp Ala Glu
 145 150 155 160
 Arg Gln Leu Leu Ser Phe Gly Asn Pro Arg Gly Thr Phe Leu Ile Arg
 165 170 175
 Glu Ser Glu Thr Thr Lys Gly Ala Tyr Ser Leu Ser Ile Arg Asp Trp
 180 185 190
 Asp Asp Met Lys Gly Asp His Val Lys His Tyr Lys Ile Arg Lys Leu
 195 200 205
 Asp Asn Gly Gly Tyr Tyr Ile Thr Thr Arg Ala Gln Phe Glu Thr Leu
 210 215 220
 Gln Gln Leu Val Gln His Tyr Ser Glu Arg Ala Ala Gly Leu Cys Cys
 225 230 235 240
 Arg Leu Val Val Pro Cys His Lys Gly Met Pro Arg Leu Thr Asp Leu
 245 250 255
 Ser Val Lys Thr Lys Asp Val Trp Glu Ile Pro Arg Glu Ser Leu Gln
 260 265 270
 Leu Ile Lys Arg Leu Gly Asn Gly Gln Phe Gly Glu Val Trp Met Gly
 275 280 285
 Met Leu Arg Leu Asn Tyr Ser Leu Ile Ser Phe Pro Val Trp Lys Ile
 290 295 300
 Pro Asn Thr Lys Asp Gly Arg
 305 310

<210> 122

<211> 387

<212> PRT

<213> Homo sapiens

<400> 122

Thr Met Ser Pro Glu Ser Phe Leu Glu Glu Ala Gln Ile Met Lys Lys
305 310 315 320

Leu Lys His Asp Lys Leu Val Gln Leu Tyr Ala Val Val Ser Glu Glu
325 330 335

Pro Ile Tyr Ile Val Thr Glu Tyr Met Asn Lys Gly Trp Ala Thr Pro
340 345 350

Leu Leu Ser Pro Ala His Ser Ala Leu Arg Gly Cys Leu Gly Glu Arg
355 360 365

Asn Gly Ser Phe Leu Leu Ala Thr Phe Leu Val Ser Ala Trp Val Lys
370 375 380

Tyr Ser His
385

<210> 123
<211> 516
<212> PRT
<213> Homo sapiens

<400> 123

Met Arg Leu Glu Leu Pro Ala Gly His Trp Glu Arg Pro Asp Leu Glu
1 5 10 15

Leu Leu Glu Lys Ser Thr Gln Gln Gly Arg Ala Trp Asp Leu Glu Leu
20 25 30

Leu Glu Lys Gly Ala Gly Ser Leu Pro Leu Tyr Val Trp Lys Val Ser
35 40 45

Leu Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala Leu Thr Glu Pro
50 55 60

Glu Ala Arg Tyr Tyr Leu Arg Gln Ile Val Leu Gly Cys Gln Tyr Leu
65 70 75 80

His Arg Asn Arg Val Ile His Arg Asp Leu Lys Leu Gly Asn Leu Phe
85 90 95

Leu Asn Glu Asp Leu Glu Val Lys Ile Gly Asp Phe Gly Leu Ala Thr
100 105 110

Lys Val Glu Tyr Asp Gly Glu Arg Lys Lys Thr Leu Cys Gly Thr Pro
115 120 125

Asn Tyr Ile Ala Pro Glu Val Leu Ser Lys Lys Gly His Ser Phe Glu
130 135 140

Val Asp Val Trp Ser Ile Gly Cys Ile Met Tyr Thr Leu Leu Val Gly
145 150 155 160

Lys Pro Pro Phe Glu Thr Ser Cys Leu Lys Glu Thr Tyr Leu Arg Ile

165 170 175

Lys	Lys	Asn	Glu	Tyr	Ser	Ile	Pro	Lys	His	Ile	Asn	Pro	Val	Ala	Ala
		180						185					190		
Ser	Leu	Ile	Gln	Lys	Met	Leu	Gln	Thr	Asp	Pro	Thr	Ala	Arg	Pro	Thr
		195					200					205			
Ile	Asn	Glu	Leu	Leu	Asn	Asp	Glu	Phe	Phe	Thr	Ser	Gly	Tyr	Ile	Pro
	210					215					220				
Ala	Arg	Leu	Pro	Ile	Thr	Cys	Leu	Thr	Ile	Pro	Pro	Arg	Phe	Ser	Ile
225					230					235					240
Ala	Pro	Ser	Ser	Leu	Asp	Pro	Ser	Asn	Arg	Lys	Pro	Leu	Thr	Val	Leu
				245					250					255	
Asn	Lys	Gly	Leu	Glu	Asn	Pro	Leu	Pro	Glu	Arg	Pro	Arg	Glu	Lys	Glu
			260					265					270		
Glu	Pro	Val	Val	Arg	Glu	Thr	Gly	Glu	Val	Val	Asp	Cys	His	Leu	Ser
		275					280					285			
Asp	Met	Leu	Gln	Gln	Leu	His	Ser	Val	Asn	Ala	Ser	Lys	Pro	Ser	Glu
	290					295					300				
Arg	Gly	Leu	Val	Arg	Gln	Glu	Glu	Ala	Glu	Asp	Pro	Ala	Cys	Ile	Pro
305					310					315					320
Ile	Phe	Trp	Val	Ser	Lys	Trp	Val	Asp	Tyr	Ser	Asp	Lys	Tyr	Gly	Leu
				325					330					335	
Gly	Tyr	Gln	Leu	Cys	Asp	Asn	Ser	Val	Gly	Val	Leu	Phe	Asn	Asp	Ser
			340					345					350		
Thr	Arg	Leu	Ile	Leu	Tyr	Asn	Asp	Gly	Asp	Ser	Leu	Gln	Tyr	Ile	Glu
		355					360					365			
Arg	Asp	Gly	Thr	Glu	Ser	Tyr	Leu	Thr	Val	Ser	Ser	His	Pro	Asn	Ser
	370					375					380				
Leu	Met	Lys	Lys	Ile	Thr	Leu	Leu	Lys	Tyr	Phe	Arg	Asn	Tyr	Met	Ser
385					390					395					400
Glu	His	Leu	Leu	Lys	Ala	Gly	Ala	Asn	Ile	Thr	Pro	Arg	Glu	Gly	Asp
				405				410						415	
Glu	Leu	Ala	Arg	Leu	Pro	Tyr	Leu	Arg	Thr	Trp	Phe	Arg	Thr	Arg	Ser
			420					425					430		
Ala	Ile	Ile	Leu	His	Leu	Ser	Asn	Gly	Ser	Val	Gln	Ile	Asn	Phe	Phe
		435					440					445			
Gln	Asp	His	Thr	Lys	Leu	Ile	Leu	Cys	Pro	Leu	Met	Ala	Ala	Val	Thr
	450					455					460				
Tyr	Ile	Asp	Glu	Lys	Arg	Asp	Phe	Arg	Thr	Tyr	Arg	Leu	Ser	Leu	Leu

465 470 475 480

Glu Glu Tyr Gly Cys Cys Lys Glu Leu Ala Ser Arg Leu Arg Tyr Ala
 485 490 495

Arg Thr Met Val Asp Lys Leu Leu Ser Ser Arg Ser Ala Ser Asn Arg
 500 505 510

Leu Lys Ala Ser
 515

<210> 124
 <211> 171
 <212> PRT
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(171)
 <223> "XAA" can be any amino acid

<400> 124

Met Ala Leu Leu Pro Pro Phe Leu Ala Ser His Arg Leu Glu Val Ser
 1 5 10 15

Arg Asp Ser Gly Trp Leu Gly Gln Cys Trp Leu Gln Gly Val Trp Glu
 20 25 30

Arg Xaa Pro His Ser Gly Leu Leu Tyr Pro Leu Gln His Pro Pro Ala
 35 40 45

Glu Phe Ser Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp Asp
 50 55 60

Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe His
 65 70 75 80

Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu Lys
 85 90 95

Phe Gly Ala Ser Ser Ser Gln Ala Gln Pro Arg Asp Ser Pro Met Thr
 100 105 110

Ala Lys Gly Pro Phe Cys Pro Arg Pro Cys Pro Cys Ala Gly Pro Thr
 115 120 125

Tyr Ser Pro Thr Tyr Trp Cys Pro Ala Pro Leu Gly Thr Gln Ser Pro
 130 135 140

Pro Asp Arg Pro Val Glu Glu Val Glu Glu Leu Ser Pro Gln Asn Tyr
 145 150 155 160

Trp Pro Val Val Trp Thr Pro Gly Pro His Phe
 165 170

<210> 125
<211> 134
<212> PRT
<213> Homo sapiens

<400> 125

Met Ala Leu Leu Pro Pro Phe Leu Ala Ser His Arg Leu Glu Val Ser
1 5 10 15
Arg Asp Ser Gly Trp Leu Gly Gln Cys Trp Leu Gln Gly Val Trp Glu
20 25 30
Arg Gly Leu Thr Val Ala Phe Ser Ile Leu Cys Asn Thr Leu Gln Pro
35 40 45
Glu Phe Ser Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp Asp
50 55 60
Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe His
65 70 75 80
Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu Lys
85 90 95
Phe Gly Ala Ser Ser Ser Gln Ala Gln Pro Arg Asp Ser Pro Met Thr
100 105 110
Ala Lys Gly Pro Phe Cys Pro Arg Pro Cys Pro Cys Ala Gly Pro Thr
115 120 125
Tyr Ser Pro Thr Tyr Trp
130

<210> 126
<211> 233
<212> PRT
<213> Homo sapiens

<400> 126

Met Ala Leu Leu Pro Pro Phe Leu Ala Ser His Arg Leu Glu Val Ser
1 5 10 15
Arg Asp Ser Gly Trp Leu Gly Gln Cys Trp Leu Gln Gly Val Trp Glu
20 25 30
Arg Gly Leu Thr Val Ala Phe Ser Ile Leu Cys Asn Thr Leu Gln Pro
35 40 45
Glu Phe Ser Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp Asp
50 55 60
Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe His
65 70 75 80
Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu Lys

Ala Val Ile Val Gly Lys Glu Ser Lys Ser Ala Ala Thr Lys Glu Glu
115 120 125

Ser Val Ser Leu Lys Glu Lys Thr Lys Pro Leu Thr Pro Ser Ile Gly
130 135 140

Ala Lys Glu Lys Glu Gln His Val Ala Leu Val Thr Ser Thr Leu Pro
145 150 155 160

Pro Leu Pro Leu Pro Pro Met Leu Pro Glu Asp Lys Glu Ala Asp Ser
165 170 175

Leu Arg Gly Asn Ile Ser Val Lys Ala Val Lys Lys Glu Val Glu Lys
180 185 190

Lys Leu Arg Cys Leu Leu Ala Asp Leu Pro Leu Pro Pro Glu Leu Pro
195 200 205

Gly Gly Asp Asp Leu Ser Lys Ser Pro Glu Glu Lys Lys Thr Ala Thr
210 215 220

Gln Leu His Ser Lys Arg Arg Pro Lys Tyr Val Leu Ala Phe Tyr Leu
225 230 235 240

Leu Leu Asn

<210> 128

<211> 330

<212> PRT

<213> Homo sapiens

<400> 128

Met Ser Ala Lys Val Arg Leu Lys Lys Leu Glu Gln Leu Leu Leu Asp
1 5 10 15

Gly Pro Trp Arg Asn Glu Ser Ala Leu Ser Val Glu Thr Leu Leu Asp
20 25 30

Val Leu Val Cys Leu Tyr Thr Glu Cys Ser His Ser Ala Leu Arg Arg
35 40 45

Asp Lys Tyr Val Ala Glu Phe Leu Glu Trp Ala Lys Pro Phe Thr Gln
50 55 60

Leu Val Lys Glu Met Gln Leu His Arg Glu Asp Phe Glu Ile Ile Lys
65 70 75 80

Val Ile Gly Arg Gly Ala Phe Gly Glu Val Ala Val Val Lys Met Lys
85 90 95

Asn Thr Glu Arg Ile Tyr Ala Met Lys Ile Leu Asn Lys Trp Glu Met
100 105 110

Leu Lys Arg Ala Glu Thr Ala Cys Phe Arg Glu Glu Arg Asp Val Leu
115 120 125

Val Asn Gly Asp Cys Gln Trp Ile Thr Ala Leu His Tyr Ala Phe Gln
130 135 140

Asp Glu Asn His Leu Tyr Leu Val Met Asp Tyr Tyr Val Gly Gly Asp
145 150 155 160

Leu Leu Thr Leu Leu Ser Lys Phe Glu Asp Lys Leu Pro Glu Asp Met
165 170 175

Ala Arg Phe Tyr Ile Gly Glu Met Val Leu Ala Ile Asp Ser Ile His
180 185 190

Gln Leu His Tyr Val His Arg Asp Ile Lys Pro Asp Asn Val Leu Leu
195 200 205

Asp Val Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys
210 215 220

Met Asn Asp Asp Gly Thr Val Gln Ser Ser Val Ala Val Gly Thr Pro
225 230 235 240

Asp Tyr Ile Ser Pro Glu Ile Leu Gln Ala Met Glu Asp Gly Met Gly
245 250 255

Lys Tyr Gly Pro Glu Cys Asp Trp Trp Ser Leu Gly Val Cys Met Tyr
260 265 270

Glu Met Leu Tyr Gly Glu Thr Pro Phe Tyr Ala Glu Ser Leu Val Glu
275 280 285

Thr Tyr Gly Lys Ile Met Asn His Glu Glu Arg Phe Gln Phe Pro Ser
290 295 300

His Val Thr Asp Val Ser Glu Glu Ala Lys Asp Leu Ile Gln Arg Leu
305 310 315 320

Ser Cys Ile Gln Arg Thr Pro Tyr Leu Gln
325 330

<210> 129

<211> 246

<212> PRT

<213> Homo sapiens

<400> 129

Met Ser Ala Lys Val Arg Leu Lys Lys Leu Glu Gln Leu Leu Leu Asp
1 5 10 15

Gly Pro Trp Arg Asn Glu Ser Ala Leu Ser Val Glu Thr Leu Leu Asp
20 25 30

Val Leu Val Cys Leu Tyr Thr Glu Cys Ser His Ser Ala Leu Arg Arg
35 40 45

Asp Lys Tyr Val Ala Glu Phe Leu Glu Trp Ala Lys Pro Phe Thr Gln

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Leu Lys Thr Val Leu Leu Leu Ala Asp Gln Met Ile Ser Arg Ile Glu
 65 70 75 80
 Tyr Ile His Ser Lys Asn Phe Ile His Arg Asp Val Lys Pro Asp Asn
 85 90 95
 Phe Leu Met Gly Leu Gly Lys Lys Gly Asn Leu Val Tyr Ile Ile Asp
 100 105 110
 Phe Gly Leu Ala Lys Lys Tyr Arg Asp Ala Arg Thr His Gln His Ile
 115 120 125
 Pro Tyr Arg Glu Asn Lys Asn Leu Thr Gly Thr Ala Arg Tyr Ala Ser
 130 135 140
 Ile Asn Thr His Leu Gly Ile Glu Gln Ser Arg Arg Asp Asp Leu Glu
 145 150 155 160
 Ser Leu Gly Tyr Val Leu Met Tyr Phe Asn Leu Gly Ser Leu Pro Trp
 165 170 175
 Gln Gly Leu Lys Ala Ala Thr Lys Arg Gln Lys Tyr Glu Arg Ile Ser
 180 185 190
 Glu Lys Lys Met Ser Thr Pro Ile Glu Val Leu Cys Lys Gly Tyr Pro
 195 200 205
 Ser Glu Phe Ala Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp
 210 215 220
 Asp Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe
 225 230 235 240
 His Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu
 245 250 255
 Lys Phe Gly Ala Ser Arg Ala Ala Asp Asp Ala Glu Arg Asp Ala Gly
 260 265 270
 Asp Arg Glu Glu Arg Leu Arg His Ser Arg Asn Pro Ala Thr Arg Gly
 275 280 285
 Leu Pro Ser Thr Ala Ser Gly Arg Leu Arg Gly Arg Arg Lys Val Ala
 290 295 300
 Pro Pro Thr Pro Leu Thr Pro Thr Ser His Thr Ala Asn Thr Ser Pro
 305 310 315 320
 Arg Pro Val Ser Gly Met Glu Arg Glu Arg Lys Val Ser Met Arg Leu
 325 330 335
 His Arg Gly Ala Pro Val Asn Ile Ser Ser Ser Asp Leu Thr Gly Arg
 340 345 350
 Gln Asp Thr Ser Arg Met Ser Thr Ser Gln Ile Pro Gly Arg Val Ala
 355 360 365

Val Met Leu Asp Leu Ala Lys Arg Ser Arg Ser Gly Lys Phe Arg Leu
260 265 270

Val Thr Lys Phe Lys Lys Glu Lys Asn Asn Lys Asn Lys Glu Ala His
275 280 285

Ser Ser Leu Gly Ala Pro Val His Leu Trp Gly Thr Glu Glu Val Ala
290 295 300

Ala Trp Leu Glu His Leu Ser Leu Cys Glu Tyr Lys Asp Ile Phe Thr
305 310 315 320

Arg His Asp Ile Arg Gly Ser Glu Leu Leu His Leu Glu Arg Arg Asp
325 330 335

Leu Lys Asp Leu Gly Val Thr Lys Val Gly His Met Lys Arg Ile Leu
340 345 350

Cys Gly Ile Lys Glu Leu Ser Arg Ser Ala Pro Ala Val Glu Ala Gln
355 360 365

Pro Leu Ser Ser Gln Pro Val Ala Ser Thr Ser Pro Pro Pro Arg Pro
370 375 380

Ser Leu Arg Pro Leu Ser Leu Trp Pro Leu Arg Leu Leu Pro Leu Arg
385 390 395 400

Pro Trp Ala Asp Ala Ala Ala Arg Pro Leu Leu Met Val Leu Leu Pro
405 410 415

Leu Ser Ala Thr Glu Ser Leu Arg Asp Thr Val His Gln Ser Ser Gly
420 425 430

Val Ser Asn Ile Thr Thr Gln Leu Pro Leu Lys Gln His Phe Leu Gln
435 440 445

Leu Arg Val Thr Trp Gly Thr Cys Val Thr Ala Thr Gln Leu Ser Pro
450 455 460

Ala Cys Ala Val Gly Gln Gly Ile Gln Arg Arg Leu Ala Ser Trp Ala
465 470 475 480

Leu Leu Ala Trp Pro Arg Ala Trp Ile Val Pro Gly Ala Pro Leu Arg
485 490 495

Val Ser Phe Cys Gly Arg Thr Val Trp Leu Arg Leu Leu Ala Pro Ser
500 505 510

Gln Phe Ser Glu Thr Trp Leu Gly Pro Ser Thr Ala Ala Cys Lys Gly
515 520 525

Pro Cys Leu Leu Met Gln Leu Leu Leu Asn Lys Asn Arg Ala Leu Ser
530 535 540

Trp Phe Glu Ser Ser Met Asp Val Ser Ser Leu Val Asp Cys Asn Leu
545 550 555 560

Thr

<210> 132
<211> 213
<212> PRT
<213> Homo sapiens

<400> 132

Met Ser Asp Val Ala Ile Val Lys Glu Gly Trp Leu His Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Thr Trp Arg Pro Arg Tyr Phe Leu Leu Lys Asn Asp
20 25 30

Gly Thr Phe Ile Gly Tyr Lys Glu Arg Pro Gln Asp Val Asp Gln Arg
35 40 45

Glu Ala Pro Leu Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys
50 55 60

Thr Glu Arg Pro Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp
65 70 75 80

Thr Thr Val Ile Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg
85 90 95

Glu Glu Trp Thr Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys Lys
100 105 110

Gln Glu Glu Glu Glu Met Asp Phe Arg Ser Gly Ser Pro Ser Asp Asn
115 120 125

Ser Gly Ala Glu Glu Met Glu Val Ser Leu Ala Lys Pro Lys His Arg
130 135 140

Val Ala Leu Gly Gly Arg Ala Gly Pro Ala His Val Ser Pro His Ser
145 150 155 160

Val Ser Gln Pro Pro Trp Ala Val Cys His Gln Leu Ser Val Ile Ser
165 170 175

Leu Gly Pro Trp Ala Ser Val Gln Pro Gly Gly Thr Arg Cys Asn Leu
180 185 190

Thr Met Val Cys Trp Pro Ala Pro Ser Pro Gly Gly Gly Arg His Thr
195 200 205

Ala Ala Pro Gln His
210

<210> 133
<211> 425
<212> PRT
<213> Homo sapiens

132 213 PRT Homo sapiens

<400> 133

Met	Ile	Val	His	Asp	Asp	Val	Glu	Ser	Glu	Pro	Ala	Met	Thr	Pro	Ser	
1				5					10					15		
Lys	Glu	Gly	Thr	Leu	Ile	Val	Arg	Gln	Thr	Gln	Ser	Ala	Ser	Ser	Thr	
			20					25					30			
Leu	Gln	Lys	His	Lys	Ser	Ser	Ser	Ser	Phe	Thr	Pro	Phe	Ile	Asp	Pro	
		35					40					45				
Arg	Leu	Leu	Gln	Ile	Ser	Pro	Ser	Ser	Gly	Thr	Thr	Val	Thr	Ser	Val	
	50					55					60					
Val	Gly	Phe	Ser	Cys	Asp	Gly	Met	Arg	Pro	Glu	Ala	Ile	Arg	Gln	Asp	
65					70					75					80	
Pro	Thr	Arg	Lys	Gly	Ser	Val	Val	Asn	Val	Asn	Pro	Thr	Asn	Thr	Arg	
				85					90					95		
Pro	Gln	Ser	Asp	Thr	Pro	Glu	Ile	Arg	Lys	Tyr	Lys	Lys	Arg	Phe	Asn	
			100					105					110			
Ser	Glu	Ile	Leu	Cys	Ala	Ala	Leu	Trp	Gly	Val	Asn	Leu	Leu	Val	Gly	
		115					120					125				
Thr	Glu	Ser	Gly	Leu	Met	Leu	Leu	Asp	Arg	Ser	Gly	Gln	Gly	Lys	Val	
	130					135					140					
Tyr	Pro	Leu	Ile	Asn	Arg	Arg	Arg	Phe	Gln	Gln	Met	Asp	Val	Leu	Glu	
145					150					155					160	
Gly	Leu	Asn	Val	Leu	Val	Thr	Ile	Ser	Gly	Lys	Lys	Asp	Lys	Leu	Arg	
				165					170					175		
Val	Tyr	Tyr	Leu	Ser	Trp	Leu	Arg	Asn	Lys	Ile	Leu	His	Asn	Asp	Pro	
			180					185					190			
Glu	Val	Glu	Lys	Lys	Gln	Gly	Trp	Thr	Thr	Val	Gly	Asp	Leu	Glu	Gly	
		195				200						205				
Cys	Val	His	Tyr	Lys	Val	Val	Lys	Tyr	Glu	Arg	Ile	Lys	Phe	Leu	Val	
		210				215					220					
Ile	Ala	Leu	Lys	Ser	Ser	Val	Glu	Val	Tyr	Ala	Trp	Ala	Pro	Lys	Pro	
225					230					235					240	
Tyr	His	Lys	Phe	Met	Ala	Phe	Lys	Ser	Phe	Gly	Glu	Leu	Val	His	Lys	
				245					250					255		
Pro	Leu	Leu	Val	Asp	Leu	Thr	Val	Glu	Glu	Gly	Gln	Arg	Leu	Lys	Val	
			260					265					270			
Ile	Tyr	Gly	Ser	Cys	Ala	Gly	Phe	His	Ala	Val	Asp	Val	Asp	Ser	Gly	
		275				280						285				

Ser Val Tyr Asp Ile Tyr Leu Pro Thr His Ile Gln Cys Ser Ile Lys
290 295 300

Pro His Ala Ile Ile Ile Leu Pro Asn Thr Asp Gly Met Glu Leu Leu
305 310 315 320

Val Cys Tyr Glu Asp Glu Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile
325 330 335

Thr Lys Asp Val Val Leu Gln Trp Gly Glu Met Pro Thr Ser Val Ala
340 345 350

Tyr Ile Arg Ser Asn Gln Thr Met Gly Trp Gly Glu Lys Ala Ile Glu
355 360 365

Ile Arg Ser Val Glu Thr Gly His Leu Asp Gly Val Phe Met His Lys
370 375 380

Arg Ala Gln Arg Leu Lys Phe Leu Cys Glu Arg Asn Asp Lys Val Phe
385 390 395 400

Phe Ala Ser Val Arg Ser Gly Gly Ser Ser Gln Val Tyr Phe Met Thr
405 410 415

Leu Gly Arg Thr Ser Leu Leu Ser Trp
420 425

<210> 134
<211> 515
<212> PRT
<213> Homo sapiens

<400> 134

Met Ala Ser Arg Thr Pro Arg Asn Cys Ala Val Leu Lys Gly Glu Val
1 5 10 15

Asp Leu Thr Ala Leu Ala Lys Glu Leu Arg Ala Val Glu Asp Val Arg
20 25 30

Pro Pro His Lys Val Thr Asp Tyr Ser Ser Ser Ser Glu Glu Ser Gly
35 40 45

Thr Thr Asp Glu Glu Asp Asp Asp Val Glu Gln Glu Gly Ala Asp Glu
50 55 60

Ser Thr Ser Gly Pro Glu Asp Thr Arg Ala Ala Ser Ser Leu Asn Leu
65 70 75 80

Ser Asn Gly Glu Thr Glu Ser Val Lys Thr Met Ile Val His Asp Asp
85 90 95

Val Glu Ser Glu Pro Ala Met Thr Pro Ser Lys Glu Gly Thr Leu Ile
100 105 110

Val Arg Gln Thr Gln Ser Ala Ser Ser Thr Leu Gln Lys His Lys Ser
115 120 125

134 515 PRT Homo sapiens 134

Ser Ser Ser Phe Thr Pro Phe Ile Asp Pro Arg Leu Leu Gln Ile Ser
 130 135 140

Pro Ser Ser Gly Thr Thr Val Thr Ser Val Val Gly Phe Ser Cys Asp
 145 150 155 160

Gly Met Arg Pro Glu Ala Ile Arg Gln Asp Pro Thr Arg Lys Gly Ser
 165 170 175

Val Val Asn Val Asn Pro Thr Asn Thr Arg Pro Gln Ser Asp Thr Pro
 180 185 190

Glu Ile Arg Lys Tyr Lys Lys Arg Phe Asn Ser Glu Ile Leu Cys Ala
 195 200 205

Ala Leu Trp Gly Val Asn Leu Leu Val Gly Thr Glu Ser Gly Leu Met
 210 215 220

Leu Leu Asp Arg Ser Gly Gln Gly Lys Val Tyr Pro Leu Ile Asn Arg
 225 230 235 240

Arg Arg Phe Gln Gln Met Asp Val Leu Glu Gly Leu Asn Val Leu Val
 245 250 255

Thr Ile Ser Gly Lys Lys Asp Lys Leu Arg Val Tyr Tyr Leu Ser Trp
 260 265 270

Leu Arg Asn Lys Ile Leu His Asn Asp Pro Glu Val Glu Lys Lys Gln
 275 280 285

Gly Trp Thr Thr Val Gly Asp Leu Glu Gly Cys Val His Tyr Lys Val
 290 295 300

Val Lys Tyr Glu Arg Ile Lys Phe Leu Val Ile Ala Leu Lys Ser Ser
 305 310 315 320

Val Glu Val Tyr Ala Trp Ala Pro Lys Pro Tyr His Lys Phe Met Ala
 325 330 335

Phe Lys Ser Phe Gly Glu Leu Val His Lys Pro Leu Leu Val Asp Leu
 340 345 350

Thr Val Glu Glu Gly Gln Arg Leu Lys Val Ile Tyr Gly Ser Cys Ala
 355 360 365

Gly Phe His Ala Val Asp Val Asp Ser Gly Ser Val Tyr Asp Ile Tyr
 370 375 380

Leu Pro Thr His Ile Gln Cys Ser Ile Lys Pro His Ala Ile Ile Ile
 385 390 395 400

Leu Pro Asn Thr Asp Gly Met Glu Leu Leu Val Cys Tyr Glu Asp Glu
 405 410 415

Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile Thr Lys Asp Val Val Leu
 420 425 430

465

<210> 136
<211> 666
<212> PRT
<213> Homo sapiens

<220>
<221> -
<222> (1)..(666)
<223> "XAA" can be any amino acid

<400> 136

Met	Asp	Cys	Gln	Leu	Ser	Ile	Leu	Leu	Leu	Leu	Ser	Cys	Ser	Val	Leu
1				5					10					15	
Asp	Ser	Phe	Gly	Glu	Leu	Ile	Pro	Gln	Pro	Ser	Asn	Glu	Val	Asn	Leu
			20					25					30		
Leu	Asp	Ser	Lys	Thr	Ile	Gln	Gly	Glu	Leu	Gly	Trp	Ile	Ser	Tyr	Pro
			35				40					45			
Ser	His	Gly	Trp	Glu	Glu	Ile	Ser	Gly	Val	Asp	Glu	His	Tyr	Thr	Pro
	50					55					60				
Ile	Arg	Thr	Tyr	Gln	Val	Cys	Asn	Val	Met	Asp	His	Ser	Gln	Asn	Asn
65					70					75					80
Trp	Leu	Arg	Thr	Asn	Trp	Val	Pro	Arg	Asn	Ser	Ala	Gln	Lys	Ile	Tyr
				85					90					95	
Val	Glu	Leu	Lys	Phe	Thr	Leu	Arg	Asp	Cys	Asn	Ser	Ile	Pro	Leu	Val
			100					105					110		
Leu	Gly	Thr	Cys	Lys	Glu	Thr	Phe	Asn	Leu	Tyr	Tyr	Met	Glu	Ser	Asp
			115				120					125			
Asp	Asp	His	Gly	Val	Lys	Phe	Arg	Glu	His	Gln	Phe	Thr	Lys	Ile	Asp
	130					135					140				
Thr	Ile	Ala	Ala	Asp	Glu	Ser	Phe	Thr	Gln	Met	Asp	Leu	Gly	Asp	Arg
145					150					155					160
Ile	Leu	Lys	Leu	Asn	Thr	Glu	Ile	Arg	Glu	Val	Gly	Pro	Val	Asn	Lys
				165					170					175	
Lys	Gly	Phe	Tyr	Leu	Ala	Phe	Gln	Asp	Val	Gly	Ala	Cys	Val	Ala	Leu
			180					185					190		
Val	Ser	Val	Arg	Val	Tyr	Phe	Lys	Lys	Cys	Pro	Phe	Thr	Val	Lys	Asn
			195				200					205			
Leu	Ala	Met	Phe	Pro	Asp	Thr	Val	Pro	Met	Asp	Ser	Gln	Ser	Leu	Val
	210					215					220				

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Glu Val Arg Gly Ser Cys Val Asn Asn Ser Lys Glu Glu Asp Pro Pro
 225 230 235 240
 Arg Met Tyr Cys Ser Thr Glu Gly Glu Trp Leu Val Pro Ile Gly Lys
 245 250 255
 Cys Ser Cys Asn Ala Gly Tyr Glu Glu Arg Gly Phe Met Cys Gln Ala
 260 265 270
 Cys Arg Pro Gly Phe Tyr Lys Ala Leu Asp Gly Asn Met Lys Cys Ala
 275 280 285
 Lys Cys Pro Pro His Ser Ser Thr Gln Glu Asp Gly Ser Met Asn Cys
 290 295 300
 Arg Cys Glu Asn Asn Tyr Phe Arg Ala Asp Lys Asp Pro Pro Ser Met
 305 310 315 320
 Ala Cys Thr Arg Pro Pro Ser Ser Pro Arg Asn Val Ile Ser Asn Ile
 325 330 335
 Asn Glu Thr Ser Val Ile Leu Asp Trp Ser Trp Pro Leu Asp Thr Gly
 340 345 350
 Gly Arg Lys Asp Val Thr Phe Asn Ile Ile Cys Lys Lys Cys Gly Trp
 355 360 365
 Asn Ile Lys Gln Cys Glu Pro Cys Ser Pro Asn Val Arg Phe Leu Pro
 370 375 380
 Arg Gln Phe Gly Leu Thr Asn Thr Thr Val Thr Val Thr Asp Leu Leu
 385 390 395 400
 Ala His Thr Asn Tyr Thr Phe Glu Ile Asp Ala Val Asn Gly Val Ser
 405 410 415
 Glu Leu Ser Ser Pro Pro Arg Gln Phe Ala Ala Val Ser Ile Thr Thr
 420 425 430
 Asn Gln Ala Ala Pro Ser Pro Val Leu Thr Ile Lys Lys Asp Arg Thr
 435 440 445
 Ser Arg Asn Ser Ile Ser Leu Ser Trp Gln Glu Pro Glu His Pro Asn
 450 455 460
 Gly Ile Ile Leu Asp Tyr Glu Val Lys Tyr Tyr Glu Lys Gln Glu Gln
 465 470 475 480
 Glu Thr Ser Tyr Thr Ile Leu Arg Ala Arg Gly Thr Asn Val Thr Ile
 485 490 495
 Ser Ser Leu Lys Pro Asp Thr Ile Tyr Val Phe Gln Ile Arg Ala Arg
 500 505 510
 Thr Ala Ala Gly Tyr Gly Thr Asn Ser Arg Lys Phe Glu Phe Glu Thr
 515 520 525

Met Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys Leu
130 135 140

Met	Glu	Asp	Gly	Thr	Val	Gln	Ser	Ser	Val	Ala	Val	Gly	Thr	Pro	Asp
145					150					155					160

Tyr Ile Ser Pro Glu Ile Leu Gln Ala Met Glu Asp Gly Lys Gly Arg
165 170 175

Tyr Gly Pro Glu Cys Asp Trp Trp Ser Leu Gly Val Cys Met Tyr Glu
180 185 190

Met Leu Tyr Gly Glu Thr Pro Phe Tyr Ala Glu Ser Leu Val Glu Thr
195 200 205

Tyr Gly Lys Ile Met Asn His Lys Glu Arg Phe Gln Phe Pro Ala Gln
210 215 220

Val	Thr	Asp	Val	Ser	Glu	Asn	Ala	Lys	Asp	Leu	Ile	Arg	Arg	Leu	Ile
225					230					235					240

Cys Ser Arg Glu His Arg Leu Gly Gln Asn Gly Ile Glu Asp Phe Lys
245 250 255

Lys His Pro Phe Phe Ser Gly Ile Asp Trp Asp Asn Ile Arg Asn Cys
260 265 270

Glu Ala Pro Tyr Ile Pro Glu Val Ser Ser Pro Thr Asp Thr Ser Asn
275 280 285

Phe Asp Val Asp Asp Asp Cys Leu Lys Asn Ser Glu Thr Met Pro Pro
290 295 300

Pro Thr His Thr Ala Phe Ser Gly His His Leu Pro Phe Val Gly Phe
305 310 315 320

Thr Tyr Thr Ser Ser Cys Val Leu Ser Asp Arg Ser Cys Leu Arg Val
325 330 335

Thr Ala Gly Pro Thr Ser Leu Asp Leu Asp Val Asn Val Gln Arg Thr
340 345 350

Leu Asp Asn Asn Leu Ala Thr Glu Ala Tyr Glu Arg Arg Ile Lys Arg
355 360 365

Leu Glu Gln Glu Lys Leu Glu Leu Ser Arg Lys Leu Gln Glu Ser Thr
370 375 380

Gln Thr Val Gln Ala Leu Gln Tyr Ser Thr Val Asp Gly Pro Leu Thr
385 390 395 400

Ala Ser Lys Asp Leu Glu Ile Lys Asn Leu Lys Glu Glu Ile Glu Lys
405 410 415

Leu Arg Lys Gln Val Thr Glu Ser Ser His Leu Glu Gln Gln Leu Glu
420 425 430

225		230		235		240
Gln Gly Ala Gln Arg Arg Pro Ala Ala Pro Pro Gln Thr Ser Trp Arg						
		245		250		255

Val Trp Arg Pro Gly Ser
260

<210> 139
<211> 203
<212> PRT
<213> Homo sapiens

<400> 139

Met Glu Val Val Asp Pro Gln Gln Leu Gly Met Phe Thr Glu Gly Glu
1 5 10 15

Leu Met Ser Val Gly Met Asp Thr Phe Ile His Arg Ile Asp Ser Thr
20 25 30

Glu Val Ile Tyr Gln Pro Arg Arg Lys Arg Ala Lys Leu Ile Gly Lys
35 40 45

Tyr Leu Met Gly Asp Leu Leu Gly Glu Gly Ser Tyr Gly Lys Val Lys
50 55 60

Glu Val Leu Asp Ser Glu Thr Leu Cys Arg Arg Ala Val Lys Ile Leu
65 70 75 80

Lys Lys Lys Lys Leu Arg Arg Ile Pro Asn Gly Glu Ala Asn Val Lys
85 90 95

Lys Glu Ile Gln Leu Leu Arg Arg Leu Arg His Lys Asn Val Ile Gln
100 105 110

Leu Val Asp Val Leu Tyr Asn Glu Glu Lys Gln Lys Met Tyr Met Val
115 120 125

Met Glu Tyr Cys Val Cys Gly Met Gln Glu Met Leu Asp Ser Val Pro
130 135 140

Glu Lys Arg Phe Pro Val Cys Gln Ala His Gly Tyr Phe Cys Gln Leu
145 150 155 160

Ile Asp Gly Leu Glu Tyr Leu His Ser Gln Gly Ile Val His Lys Asp
165 170 175

Ile Lys Pro Gly Asn Leu Leu Leu Thr Thr Gly Gly Thr Leu Lys Ile
180 185 190

Ser Asp Leu Gly Val Ala Glu Val Gly Thr Cys
195 200

<210> 140
<211> 244
<212> PRT

<213> Homo sapiens

<400> 140

Met Asp Arg Glu Thr Thr Pro Leu Gly Leu Leu Trp Leu Ile Gln Val
1 5 10 15

Ile Pro Ser Lys Leu Leu Pro Ser Leu Gln Val Lys Asp Phe Leu Ser
20 25 30

Gln Leu Arg Ser Ser Asn Arg Arg Phe Ser Ile Pro Glu Ser Gly Gln
35 40 45

Gly Gly Thr Glu Met Asp Gly Phe Arg Arg Thr Ile Glu Asn Gln His
50 55 60

Ser Arg Asn Asp Val Met Val Ser Glu Trp Leu Asn Lys Leu Asn Leu
65 70 75 80

Glu Glu Pro Pro Ser Ser Val Pro Lys Lys Cys Pro Ser Leu Thr Lys
85 90 95

Arg Ser Arg Ala Gln Glu Glu Gln Val Pro Gln Ala Trp Thr Ala Gly
100 105 110

Thr Ser Ser Asp Ser Met Ala Gln Pro Pro Gln Thr Pro Glu Thr Ser
115 120 125

Thr Phe Arg Asn Gln Met Pro Ser Pro Thr Ser Thr Gly Thr Pro Ser
130 135 140

Pro Gly Pro Arg Gly Asn Gln Gly Ala Glu Arg Gln Gly Met Asn Trp
145 150 155 160

Ser Cys Arg Thr Pro Glu Pro Asn Pro Val Thr Gly Arg Pro Leu Val
165 170 175

Asn Ile Tyr Asn Cys Ser Gly Val Gln Val Gly Asp Asn Asn Tyr Leu
180 185 190

Thr Met Gln Gln Thr Thr Ala Leu Pro Thr Trp Gly Leu Ala Pro Ser
195 200 205

Gly Lys Gly Arg Gly Leu Gln His Pro Pro Pro Val Gly Ser Gln Glu
210 215 220

Gly Pro Lys Asp Pro Glu Ala Trp Ser Arg Pro Gln Gly Trp Tyr Asn
225 230 235 240

His Ser Gly Lys

<210> 141

<211> 222

<212> PRT

<213> Homo sapiens

<400> 141

Met Val Lys Leu Tyr Leu Tyr Gln Lys Asn Val Lys Ile Ala Ile Phe
1 5 10 15

Asp Leu Lys Ser Arg Gln Asn Phe Phe Val Tyr Phe Arg Glu Glu Gln
20 25 30

Ala Arg Glu Leu Tyr Arg Arg Leu Arg Glu Lys Pro Arg Asp Gln Arg
35 40 45

Thr Glu Gly Asp Ser Gln Glu Met Val Arg Leu Leu Leu Gln Ala Ile
50 55 60

Gln Ser Phe Glu Lys Lys Val Arg Val Ile Tyr Thr Gln Leu Ser Lys
65 70 75 80

Thr Val Val Cys Lys Gln Lys Ala Leu Glu Leu Leu Pro Lys Val Glu
85 90 95

Glu Val Val Ser Leu Met Asn Glu Asp Glu Lys Thr Val Val Arg Leu
100 105 110

Gln Glu Lys Arg Gln Lys Glu Leu Trp Asn Leu Leu Lys Ile Ala Cys
115 120 125

Ser Lys Val Arg Gly Pro Val Ser Gly Ser Pro Asp Ser Met Asn Ala
130 135 140

Ser Arg Leu Ser Gln Pro Gly Gln Leu Met Ser Gln Pro Ser Thr Ala
145 150 155 160

Ser Asn Ser Leu Pro Glu Pro Ala Lys Lys Ser Glu Glu Leu Val Ala
165 170 175

Glu Ala His Asn Leu Cys Thr Leu Leu Glu Asn Ala Ile Gln Asp Thr
180 185 190

Val Arg Glu Gln Asp Gln Ser Phe Thr Ala Leu Asp Trp Ser Trp Leu
195 200 205

Gln Thr Glu Glu Glu Glu His Ser Cys Leu Glu Gln Ala Ser
210 215 220

<210> 142

<211> 409

<212> PRT

<213> Homo sapiens

<400> 142

Met Arg Leu Thr Leu Leu Cys Cys Thr Trp Arg Glu Glu Arg Met Gly
1 5 10 15

Glu Glu Gly Ser Glu Leu Pro Val Cys Ala Ser Cys Gly Gln Arg Ile
20 25 30

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Tyr	Asp	Gly	Gln	Tyr	Leu	Gln	Ala	Leu	Asn	Ala	Asp	Trp	His	Ala	Asp	
	35						40					45				
Cys	Phe	Arg	Cys	Cys	Asp	Cys	Ser	Ala	Ser	Leu	Ser	His	Gln	Tyr	Tyr	
	50					55					60					
Glu	Lys	Asp	Gly	Gln	Leu	Phe	Cys	Lys	Lys	Asp	Tyr	Trp	Ala	Arg	Tyr	
65					70					75					80	
Gly	Glu	Ser	Cys	His	Gly	Cys	Ser	Glu	Gln	Ile	Thr	Lys	Gly	Leu	Val	
				85					90					95		
Met	Val	Ala	Gly	Glu	Leu	Lys	Tyr	His	Pro	Glu	Cys	Phe	Ile	Cys	Leu	
		100						105					110			
Thr	Cys	Gly	Thr	Phe	Ile	Gly	Asp	Gly	Asp	Thr	Tyr	Thr	Leu	Val	Glu	
		115					120						125			
His	Ser	Lys	Leu	Tyr	Cys	Gly	His	Cys	Tyr	Tyr	Gln	Thr	Val	Val	Thr	
	130					135					140					
Pro	Val	Ile	Glu	Gln	Ile	Leu	Pro	Asp	Ser	Pro	Gly	Ser	His	Leu	Pro	
145					150					155					160	
His	Thr	Val	Thr	Leu	Val	Ser	Ile	Pro	Ala	Ser	Ser	His	Gly	Lys	Arg	
				165					170					175		
Gly	Leu	Ser	Val	Ser	Ile	Asp	Pro	Pro	His	Gly	Pro	Pro	Gly	Cys	Gly	
			180					185					190			
Thr	Glu	His	Ser	His	Thr	Val	Arg	Val	Gln	Gly	Val	Asp	Pro	Gly	Cys	
		195					200					205				
Met	Ser	Pro	Asp	Val	Lys	Asn	Ser	Ile	His	Val	Gly	Asp	Arg	Ile	Leu	
	210					215					220					
Glu	Ile	Asn	Gly	Thr	Pro	Ile	Arg	Asn	Val	Pro	Leu	Asp	Glu	Ile	Asp	
225					230					235					240	
Leu	Leu	Ile	Gln	Glu	Thr	Ser	Arg	Leu	Leu	Gln	Leu	Thr	Leu	Glu	His	
			245						250					255		
Asp	Pro	His	Asp	Thr	Leu	Gly	His	Gly	Leu	Gly	Pro	Glu	Thr	Ser	Pro	
		260						265					270			
Leu	Ser	Ser	Pro	Ala	Tyr	Thr	Pro	Ser	Gly	Glu	Ala	Gly	Ser	Ser	Ala	
		275					280					285				
Arg	Gln	Lys	Pro	Val	Leu	Arg	Ser	Cys	Ser	Ile	Asp	Arg	Ser	Pro	Gly	
	290					295					300					
Ala	Gly	Ser	Leu	Gly	Ser	Pro	Ala	Ser	Gln	Arg	Lys	Asp	Leu	Gly	Arg	
305					310					315					320	
Ser	Glu	Ser	Leu	Arg	Val	Val	Cys	Arg	Pro	His	Arg	Ile	Phe	Arg	Pro	
				325					330					335		

Ser Asp Leu Ile His Gly Glu Val Leu Gly Lys Gly Cys Phe Gly Gln
 340 345 350

Ala Ile Lys Val Gln Ser Met Pro Gly Ser Gln Leu Asp Ser Leu Gly
 355 360 365

Gly Thr Pro Pro Ser Ser Phe Leu Pro Ser Leu Trp Lys His Ser Gly
 370 375 380

Arg Gly Ile Trp Leu Ser Asp Ser Leu Ala Ser Ala Leu Ser Ser Leu
 385 390 395 400

Gly Leu Leu Glu Leu Ile Arg Asn Arg
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<210> 143

<211> 305

<212> PRT

<213> Homo sapiens

<400> 143

Met Arg Leu Thr Leu Leu Cys Cys Thr Trp Arg Glu Glu Arg Met Gly
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Glu Glu Gly Ser Glu Leu Pro Val Cys Ala Ser Cys Gly Gln Arg Ile
 20 25 30

Tyr Asp Gly Gln Tyr Leu Gln Ala Leu Asn Ala Asp Trp His Ala Asp
 35 40 45

Cys Phe Arg Cys Cys Asp Cys Ser Ala Ser Leu Ser His Gln Tyr Tyr
 50 55 60

Glu Lys Asp Gly Gln Leu Phe Cys Lys Lys Asp Tyr Trp Ala Arg Tyr
 65 70 75 80

Gly Glu Ser Cys His Gly Cys Ser Glu Gln Ile Thr Lys Gly Leu Val
 85 90 95

Met Val Ala Gly Glu Leu Lys Tyr His Pro Glu Cys Phe Ile Cys Leu
 100 105 110

Thr Cys Gly Thr Phe Ile Gly Asp Gly Asp Thr Tyr Thr Leu Val Glu
 115 120 125

His Ser Lys Leu Tyr Cys Gly His Cys Tyr Tyr Gln Thr Val Val Thr
 130 135 140

Pro Val Ile Glu Gln Ile Leu Pro Asp Ser Pro Gly Ser His Leu Pro
 145 150 155 160

His Thr Val Thr Leu Val Ser Ile Pro Ala Ser Ser His Gly Lys Arg
 165 170 175

Gly Leu Ser Val Ser Ile Asp Pro Pro His Gly Pro Pro Gly Cys Gly
 180 185 190

Thr Glu His Ser His Thr Val Arg Val Gln Gly Val Asp Pro Gly Cys
195 200 205

Met Ser Pro Asp Val Lys Asn Ser Ile His Val Gly Asp Arg Ile Leu
210 215 220

Glu Ile Asn Gly Thr Pro Ile Arg Asn Val Pro Leu Asp Glu Ile Asp
225 230 235 240

Leu Leu Ile Gln Glu Thr Ser Arg Leu Leu Gln Leu Thr Leu Glu His
245 250 255

Asp Pro His Asp Thr Leu Gly His Gly Leu Gly Pro Glu Thr Ser Pro
260 265 270

Leu Ser Ser Pro Ala Tyr Thr Pro Ser Gly Glu Ala Gly Ser Ser Ala
275 280 285

Arg Gln Lys Pro Val Phe Ala Arg Thr Trp Val Ala Leu Ser Pro Ser
290 295 300

Ala
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<210> 144
<211> 780
<212> PRT
<213> Homo sapiens

<400> 144

Met Ala Ser Asp Ala Val Gln Ser Glu Pro Arg Ser Trp Ser Leu Leu
1 5 10 15

Glu Gln Leu Gly Leu Ala Gly Ala Asp Leu Ala Ala Pro Gly Val Gln
20 25 30

Gln Gln Leu Glu Leu Glu Arg Glu Arg Leu Arg Arg Glu Ile Arg Lys
35 40 45

Glu Leu Lys Leu Lys Glu Gly Ala Glu Asn Leu Arg Arg Ala Thr Thr
50 55 60

Asp Leu Gly Arg Ser Leu Gly Pro Val Glu Leu Leu Leu Arg Gly Ser
65 70 75 80

Ser Arg Arg Leu Asp Leu Leu His Gln Gln Leu Gln Glu Leu His Ala
85 90 95

His Val Val Leu Pro Asp Pro Ala Ala Thr His Asp Gly Pro Gln Ser
100 105 110

Pro Gly Ala Gly Gly Pro Thr Cys Ser Ala Thr Asn Leu Ser Arg Val
115 120 125

Ala Gly Leu Glu Lys Gln Leu Ala Ile Glu Leu Lys Val Lys Gln Gly

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130	135	140
Ala Glu Asn Met Ile Gln Thr Tyr Ser Asn Gly Ser Thr Lys Asp Arg		
145	150	155 160
Lys Leu Leu Leu Thr Ala Gln Gln Met Leu Gln Asp Ser Lys Thr Lys		
	165	170 175
Ile Asp Ile Ile Arg Met Gln Leu Arg Arg Ala Leu Gln Ala Asp Gln		
	180	185 190
Leu Glu Asn Gln Ala Ala Pro Asp Asp Thr Gln Gly Ser Pro Asp Leu		
	195	200 205
Gly Ala Val Glu Leu Arg Ile Glu Glu Leu Arg His His Phe Arg Val		
	210	215 220
Glu His Ala Val Ala Glu Gly Ala Lys Asn Val Leu Arg Leu Leu Ser		
	225	230 235 240
Ala Ala Lys Ala Pro Asp Arg Lys Ala Val Ser Glu Ala Gln Glu Lys		
	245	250 255
Leu Thr Glu Ser Asn Gln Lys Leu Gly Leu Leu Arg Glu Ala Leu Glu		
	260	265 270
Arg Arg Leu Gly Glu Leu Pro Ala Asp His Pro Lys Gly Arg Leu Leu		
	275	280 285
Arg Glu Glu Leu Ala Ala Ala Ser Ser Ala Ala Phe Ser Thr Arg Leu		
	290	295 300
Ala Gly Pro Phe Pro Ala Thr His Tyr Ser Thr Leu Cys Lys Pro Ala		
	305	310 315 320
Pro Leu Thr Gly Thr Leu Glu Val Arg Val Val Gly Cys Arg Asp Leu		
	325	330 335
Pro Glu Thr Ile Pro Trp Asn Pro Thr Pro Ser Met Gly Gly Pro Gly		
	340	345 350
Thr Pro Asp Ser Arg Pro Pro Phe Leu Ser Arg Pro Ala Arg Gly Leu		
	355	360 365
Tyr Ser Arg Ser Gly Ser Leu Ser Gly Arg Ser Ser Leu Lys Ala Glu		
	370	375 380
Ala Glu Asn Thr Ser Glu Val Ser Thr Val Leu Lys Leu Asp Asn Thr		
	385	390 395 400
Val Val Gly Gln Thr Ser Trp Lys Pro Cys Gly Pro Asn Ala Trp Asp		
	405	410 415
Gln Ser Phe Thr Leu Glu Leu Glu Arg Ala Arg Glu Leu Glu Leu Ala		
	420	425 430
Val Phe Trp Arg Asp Gln Arg Gly Leu Cys Ala Leu Lys Phe Leu Lys		

435					440					445					
Leu	Glu	Asp	Phe	Leu	Asp	Asn	Glu	Arg	His	Glu	Val	Gln	Leu	Asp	Met
450						455					460				
Glu	Pro	Gln	Gly	Cys	Leu	Val	Ala	Glu	Val	Thr	Phe	Arg	Asn	Pro	Val
465					470					475					480
Ile	Glu	Arg	Ile	Pro	Arg	Leu	Arg	Arg	Gln	Lys	Lys	Ile	Phe	Ser	Lys
				485					490					495	
Gln	Gln	Gly	Lys	Ala	Phe	Gln	Arg	Ala	Arg	Gln	Met	Asn	Ile	Asp	Val
			500					505					510		
Ala	Thr	Trp	Val	Arg	Leu	Leu	Arg	Arg	Leu	Ile	Pro	Asn	Ala	Thr	Gly
			515				520					525			
Thr	Gly	Thr	Phe	Ser	Pro	Gly	Ala	Ser	Pro	Gly	Ser	Glu	Ala	Arg	Thr
	530					535					540				
Thr	Gly	Asp	Ile	Ser	Val	Glu	Lys	Leu	Asn	Leu	Gly	Thr	Asp	Ser	Asp
545					550					555					560
Ser	Ser	Pro	Gln	Lys	Ser	Ser	Arg	Asp	Pro	Pro	Ser	Ser	Pro	Ser	Ser
				565					570					575	
Leu	Ser	Ser	Pro	Ile	Gln	Glu	Ser	Thr	Ala	Pro	Glu	Leu	Pro	Ser	Glu
			580					585					590		
Thr	Gln	Glu	Thr	Pro	Gly	Pro	Ala	Leu	Cys	Ser	Pro	Leu	Arg	Lys	Ser
			595				600					605			
Pro	Leu	Thr	Leu	Glu	Asp	Phe	Lys	Phe	Leu	Ala	Val	Leu	Gly	Arg	Gly
	610					615					620				
His	Phe	Gly	Lys	Val	Leu	Leu	Ser	Glu	Phe	Arg	Pro	Ser	Gly	Glu	Leu
625					630					635					640
Phe	Ala	Ile	Lys	Ala	Leu	Lys	Lys	Gly	Asp	Ile	Val	Ala	Arg	Asp	Glu
				645					650					655	
Val	Glu	Ser	Leu	Met	Cys	Glu	Lys	Arg	Ile	Leu	Ala	Ala	Val	Thr	Ser
			660					665					670		
Ala	Gly	His	Pro	Phe	Leu	Val	Asn	Leu	Phe	Gly	Cys	Phe	Gln	Thr	Pro
			675				680					685			
Glu	His	Val	Cys	Phe	Val	Met	Glu	Tyr	Ser	Ala	Gly	Gly	Asp	Leu	Met
	690					695					700				
Leu	His	Ile	His	Ser	Asp	Val	Phe	Ser	Glu	Pro	Arg	Ala	Ile	Phe	Tyr
705					710					715					720
Ser	Ala	Cys	Arg	Leu	Pro	Pro	Pro	Phe	Val	Pro	Thr	Leu	Ser	Gly	Arg
				725					730					735	
Thr	Asp	Val	Ser	Asn	Phe	Asp	Glu	Glu	Phe	Thr	Gly	Glu	Ala	Pro	Thr

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Glu	His	Ala	Val	Ala	Glu	Gly	Ala	Lys	Asn	Val	Leu	Arg	Leu	Leu	Ser	225		230		235		240
Ala	Ala	Lys	Ala	Pro	Asp	Arg	Lys	Ala	Val	Ser	Glu	Ala	Gln	Glu	Lys		245		250		255	
Leu	Thr	Glu	Ser	Asn	Gln	Lys	Leu	Gly	Leu	Leu	Arg	Glu	Ala	Leu	Glu		260		265		270	
Arg	Arg	Leu	Gly	Glu	Leu	Pro	Ala	Asp	His	Pro	Lys	Gly	Arg	Leu	Leu		275		280		285	
Arg	Glu	Glu	Leu	Ala	Ala	Ala	Ser	Ser	Ala	Ala	Phe	Ser	Thr	Arg	Leu		290		295		300	
Ala	Gly	Pro	Phe	Pro	Ala	Thr	His	Tyr	Ser	Thr	Leu	Cys	Lys	Pro	Ala	305		310		315		320
Pro	Leu	Thr	Gly	Thr	Leu	Glu	Val	Arg	Val	Val	Gly	Cys	Arg	Asp	Leu		325		330		335	
Pro	Glu	Thr	Ile	Pro	Trp	Asn	Pro	Thr	Pro	Ser	Met	Gly	Gly	Pro	Gly		340		345		350	
Thr	Pro	Asp	Ser	Arg	Pro	Pro	Phe	Leu	Ser	Arg	Pro	Ala	Arg	Gly	Leu		355		360		365	
Tyr	Ser	Arg	Ser	Gly	Ser	Leu	Ser	Gly	Arg	Ser	Ser	Leu	Lys	Ala	Glu	370		375		380		
Ala	Glu	Asn	Thr	Ser	Glu	Val	Ser	Thr	Val	Leu	Lys	Leu	Asp	Asn	Thr	385		390		395		400

His

<210> 146
 <211> 96
 <212> PRT
 <213> Homo sapiens

<400> 146

Met	Gln	Ser	Phe	Leu	Val	Glu	Gly	Arg	Phe	Lys	His	Glu	Met	Phe	Glu	1		5		10		15
Lys	Val	Phe	Ala	Glu	Glu	Arg	Asn	Gly	Gly	Gln	Arg	Leu	Leu	Cys	Ala		20		25		30	
Thr	Asp	Val	Pro	Ile	Arg	Thr	Val	Ser	Ser	Ala	Ala	Ser	Gln	Gly	Leu		35		40		45	
His	Met	Gln	Asn	Asp	Asp	Ala	Cys	Leu	Gly	Ala	Ala	Ser	Pro	Ser	Ala	50		55		60		
Ala	Ser	Trp	Ser	Arg	Arg	Ser	Ala	Glu	Ser	Lys	Val	Ser	Leu	Cys	Trp	65		70		75		80

Lys Leu Lys Trp Lys Glu Asp Leu Val Trp Phe Tyr Ser Gln Ser His
85 90 95

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<210> 147
<211> 333
<212> PRT
<213> Homo sapiens
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<400> 147

Met His Arg Tyr Phe Glu Ser Pro Arg Arg Leu Leu Pro Val His Phe
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Cys Cys Cys Gln Trp Arg Gly Gly Gly Val Asp Phe Glu Cys Leu Leu
20 25 30

Gly Gly Val Trp Asp Arg Cys Arg Lys Val Leu Arg Ala Gln Glu Cys
35 40 45

Glu Trp Pro Arg His Leu Pro Ser Ala Cys Leu Leu Ser Ser Ala Cys
50 55 60

Arg Gly Gln Pro Glu Arg Arg Ala Ala Val Val Gly Ala Gln Asp Pro
65 70 75 80

Thr Glu Pro Pro Arg Leu Ser Arg Ser Leu Ser Gly Ala Ser Pro Phe
85 90 95

Leu Gly Glu Thr Lys Gln Glu Thr Leu Thr Asn Ile Ser Ala Val Asn
100 105 110

Tyr Asp Phe Asp Glu Glu Tyr Phe Ser Asn Thr Ser Glu Leu Ala Lys
115 120 125

Asp Phe Ile Arg Arg Leu Leu Val Lys Asp Pro Lys Arg Arg Met Thr
130 135 140

Ile Ala Gln Ser Leu Glu His Ser Trp Ile Lys Ala Ile Arg Arg Arg
145 150 155 160

Asn Val Arg Gly Glu Asp Ser Gly Arg Lys Pro Glu Arg Arg Arg Leu
165 170 175

Lys Thr Thr Arg Leu Lys Glu Tyr Thr Ile Lys Ser His Ser Ser Leu
180 185 190

Pro Pro Asn Asn Ser Tyr Ala Asp Phe Glu Arg Phe Ser Lys Val Leu
195 200 205

Glu Glu Ala Ala Ala Ala Glu Glu Gly Leu Arg Glu Leu Gln Arg Ser
210 215 220

Arg Arg Leu Cys His Glu Asp Val Glu Ala Leu Ala Ala Ile Tyr Glu
225 230 235 240

Glu Lys Glu Ala Trp Tyr Arg Glu Glu Ser Asp Ser Leu Gly Gln Asp

[illegible]

	245		250		255
Leu Arg Arg	Leu Arg Gln Glu Leu	Leu Lys Thr Glu Ala	Leu Lys Arg		
260		265		270	
Gln Ala Gln Glu Glu Ala Lys Gly Ala Leu Leu Gly Thr Ser Gly Leu					
275		280		285	
Lys Arg Arg Phe Ser Arg Leu Glu Asn Arg Tyr Glu Ala Leu Ala Lys					
290		295		300	
Gln Val Ala Ser Glu Met Arg Phe Val Gln Asp Leu Val Arg Ala Leu					
305		310		315	320
Glu Gln Glu Lys Leu Gln Gly Val Glu Cys Gly Leu Arg					
	325		330		

<210> 148
 <211> 131
 <212> PRT
 <213> Homo sapiens

<400> 148

Met Leu Lys Glu Phe Leu Glu Ile Pro Phe Pro Thr Ser Pro Glu Cys					
1	5		10		15
Thr Leu Gln Pro Lys Ser Gln Gln Pro Thr Gly Lys Glu Ala Glu Glu					
	20		25		30
His Pro Thr Ser Ala Pro Leu Thr His Ser Leu Leu Pro Pro Thr Pro					
	35		40		45
Leu Trp Val Val Ser His Phe Ile Phe Asp Phe Arg Gly Glu Thr Ala					
50		55		60	
Leu His Lys Ala Ala Cys Gln Arg Asn Arg Ala Val Cys Gln Leu Leu					
65		70		75	80
Val Asp Ala Gly Ala Ser Leu Arg Lys Thr Asp Ser Lys Gly Lys Thr					
	85		90		95
Pro Gln Glu Arg Ala Gln Gln Ala Gly Asp Pro Asp Leu Ala Ala Tyr					
	100		105		110
Leu Glu Ser Arg Gln Asn Tyr Lys Val Ile Gly His Glu Asp Leu Glu					
	115		120		125

Thr Ala Val
 130

<210> 149
 <211> 272
 <212> PRT
 <213> Homo sapiens

<400> 149

Met	Arg	Gly	Ala	Ala	Arg	Leu	Gly	Arg	Pro	Gly	Arg	Ser	Cys	Leu	Pro	1	5	10	15
Gly	Pro	Ala	Leu	Arg	Ala	Pro	Pro	Arg	Pro	Pro	Leu	Leu	Leu	Leu	Leu	20	25	30	
Ala	Leu	Leu	Pro	Leu	Leu	Pro	Ala	Pro	Gly	Ala	Ala	Ala	Ala	Pro	Ala	35	40	45	
Pro	Arg	Pro	Pro	Glu	Leu	Gln	Ser	Ala	Ser	Ala	Gly	Pro	Ser	Val	Ser	50	55	60	
Leu	Tyr	Leu	Ser	Glu	Asp	Glu	Val	Arg	Arg	Leu	Ile	Gly	Leu	Asp	Ala	65	70	75	
Glu	Leu	Tyr	Tyr	Val	Arg	Asn	Asp	Leu	Ile	Ser	His	Tyr	Ala	Leu	Ser	85	90	95	
Phe	Ser	Leu	Leu	Val	Pro	Ser	Glu	Thr	Asn	Phe	Leu	His	Phe	Thr	Trp	100	105	110	
His	Ala	Lys	Ser	Lys	Val	Glu	Tyr	Lys	Leu	Gly	Phe	Gln	Val	Asp	Asn	115	120	125	
Val	Leu	Ala	Met	Asp	Met	Pro	Gln	Val	Asn	Ile	Ser	Val	Gln	Gly	Glu	130	135	140	
Val	Pro	Arg	Thr	Leu	Ser	Val	Phe	Arg	Val	Glu	Leu	Ser	Cys	Thr	Gly	145	150	155	
Lys	Val	Asp	Ser	Glu	Val	Met	Ile	Leu	Met	Gln	Leu	Asn	Leu	Thr	Val	165	170	175	
Asn	Ser	Ser	Lys	Asn	Phe	Thr	Val	Leu	Asn	Phe	Lys	Arg	Arg	Lys	Met	180	185	190	
Cys	Tyr	Lys	Lys	Leu	Glu	Glu	Val	Lys	Thr	Ser	Ala	Leu	Asp	Lys	Asn	195	200	205	
Thr	Ser	Arg	Thr	Ile	Tyr	Asp	Pro	Val	His	Ala	Ala	Pro	Thr	Thr	Ser	210	215	220	
Thr	Arg	Val	Phe	Tyr	Ile	Ser	Val	Gly	Val	Cys	Cys	Ala	Val	Ile	Phe	225	230	235	
Leu	Val	Ala	Ile	Ile	Leu	Ala	Val	Leu	His	Leu	His	Ser	Met	Lys	Arg	245	250	255	
Ile	Glu	Leu	Asp	Asp	Arg	Tyr	Cys	Thr	Tyr	Phe	Gly	Lys	Glu	Lys	Lys	260	265	270	

<210> 150

<211> 344

<212> PRT

<213> Homo sapiens

<400> 150

Met Pro Gln Val Asn Ile Ser Val Gln Gly Glu Val Pro Arg Thr Leu
1 5 10 15

Ser Val Phe Arg Val Glu Leu Ser Cys Thr Gly Lys Val Asp Ser Glu
20 25 30

Val Met Ile Leu Met Gln Leu Asn Leu Thr Val Asn Ser Ser Lys Asn
35 40 45

Phe Thr Val Leu Asn Phe Lys Arg Arg Lys Met Cys Tyr Lys Lys Leu
50 55 60

Glu Glu Val Lys Thr Ser Ala Leu Asp Lys Asn Thr Ser Arg Thr Ile
65 70 75 80

Tyr Asp Pro Val His Ala Ala Pro Thr Thr Ser Thr Arg Val Phe Tyr
85 90 95

Ile Ser Val Gly Val Cys Cys Ala Val Ile Phe Leu Val Ala Ile Ile
100 105 110

Leu Ala Val Leu His Leu His Ser Met Lys Arg Ile Glu Leu Asp Asp
115 120 125

Ser Ile Ser Ala Ser Ser Ser Ser Gln Gly Leu Ser Gln Pro Ser Thr
130 135 140

Gln Thr Thr Gln Tyr Leu Arg Ala Asp Thr Pro Asn Asn Ala Thr Pro
145 150 155 160

Ile Thr Ser Ser Tyr Tyr Pro Thr Leu Arg Ile Glu Lys Asn Asp Leu
165 170 175

Arg Ser Val Thr Leu Leu Glu Ala Lys Gly Lys Val Lys Asp Ile Ala
180 185 190

Ile Ser Arg Glu Arg Ile Thr Leu Lys Asp Val Leu Gln Glu Gly Thr
195 200 205

Phe Gly Arg Ile Phe His Gly Ile Leu Ile Asp Glu Lys Asp Pro Asn
210 215 220

Lys Glu Lys Gln Ala Phe Val Lys Thr Val Lys Asp Gln Ala Ser Glu
225 230 235 240

Ile Gln Val Thr Met Met Leu Thr Glu Ser Cys Lys Leu Arg Gly Leu
245 250 255

His His Arg Asn Leu Leu Pro Ile Thr His Val Cys Ile Glu Glu Gly
260 265 270

Glu Lys Pro Met Val Ile Leu Pro Tyr Met Asn Trp Gly Asn Leu Lys
275 280 285

Leu Phe Leu Arg Gln Cys Lys Leu Val Glu Ala Asn Asn Pro Gln Ala

290 295 300

Ile Ser Gln Gln Asp Leu Val His Met Ala Ile Gln Ile Ala Cys Gly
 305 310 315 320

Met Ser Tyr Leu Ala Arg Arg Glu Val Ile His Lys Asp Leu Ala Ala
 325 330 335

Arg Asn Cys Val Gly Pro Leu Glu
 340

<210> 151
 <211> 141
 <212> PRT
 <213> Homo sapiens

<400> 151

Met Glu Ala Ile Arg Thr Asp Asn Gln Asn Phe Ala Ser Gln Leu Arg
 1 5 10 15

Glu Ala Glu Ala Arg Asn Arg Asp Leu Glu Ala His Val Arg Gln Leu
 20 25 30

Gln Glu Arg Met Glu Leu Leu Gln Ala Glu Gly Ala Thr Ala Val Thr
 35 40 45

Gly Val Pro Ser Pro Arg Ala Thr Asp Pro Pro Ser His Leu Asp Gly
 50 55 60

Pro Pro Ala Val Ala Val Gly Gln Cys Pro Leu Val Gly Pro Gly Pro
 65 70 75 80

Met His Arg Arg His Leu Leu Leu Pro Ala Arg Val Pro Arg Pro Gly
 85 90 95

Leu Ser Glu Ala Leu Ser Leu Leu Leu Phe Ala Val Val Leu Ser Arg
 100 105 110

Ala Ala Ala Leu Gly Cys Ile Gly Leu Val Ala His Ala Gly Gln Leu
 115 120 125

Thr Ala Val Trp Arg Arg Pro Gly Ala Ala Arg Ala Pro
 130 135 140

<210> 152
 <211> 106
 <212> PRT
 <213> Homo sapiens

<400> 152

Met Glu Leu Leu Gln Ala Glu Gly Ala Thr Ala Val Thr Gly Val Pro
 1 5 10 15

Ser Pro Arg Ala Thr Asp Pro Pro Ser His Leu Asp Gly Pro Pro Ala
 20 25 30

Val Ala Val Gly Gln Cys Pro Leu Val Gly Pro Gly Pro Met His Arg
35 40 45

Arg His Leu Leu Leu Pro Ala Arg Val Pro Arg Pro Gly Leu Ser Glu
50 55 60

Ala Leu Ser Leu Leu Leu Phe Ala Val Val Leu Ser Arg Ala Ala Ala
65 70 75 80

Leu Gly Cys Ile Gly Leu Val Ala His Ala Gly Gln Leu Thr Ala Val
85 90 95

Trp Arg Arg Pro Gly Ala Ala Arg Ala Pro
100 105

<210> 153
<211> 50
<212> PRT
<213> Homo sapiens

<400> 153

Met Val Asn Leu Ser His Glu Asp Phe Glu Phe Ile Ser Gly Thr Arg
1 5 10 15

Met Arg Lys Leu Ala Arg Glu Gly Gln Lys Pro Pro Glu Gly Phe Met
20 25 30

Ala Pro Lys Ala Trp Thr Val Leu Thr Glu Tyr Tyr Lys Ser Leu Glu
35 40 45

Lys Ala
50

<210> 154
<211> 238
<212> PRT
<213> Homo sapiens

<400> 154

Met Ala Arg Thr Thr Ser Gln Leu Tyr Asp Ala Val Pro Ile Gln Ser
1 5 10 15

Ser Val Val Leu Cys Ser Cys Pro Ser Pro Ser Met Val Arg Thr Gln
20 25 30

Thr Glu Ser Ser Thr Pro Pro Gly Ile Pro Gly Gly Ser Arg Gln Gly
35 40 45

Pro Ala Met Asp Gly Thr Ala Ala Glu Pro Arg Pro Gly Ala Gly Ser
50 55 60

Leu Gln His Ala Gln Pro Pro Gln Pro Arg Lys Lys Arg Pro Glu
65 70 75 80

<400> 156

Met Ser Asp Val Thr Ile Val Lys Glu Gly Trp Val Gln Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Asn Trp Arg Pro Arg Tyr Phe Leu Leu Lys Thr Asp
20 25 30

Gly Ser Phe Ile Gly Tyr Lys Glu Lys Pro Gln Asp Val Asp Leu Pro
35 40 45

Tyr Pro Leu Asn Asn Phe Ser Val Ala Lys Cys Gln Leu Met Lys Thr
50 55 60

Glu Arg Pro Lys Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr
65 70 75 80

Thr Val Ile Glu Arg Thr Phe His Val Asp Thr Pro Glu Glu Arg Glu
85 90 95

Glu Trp Thr Glu Ala Ile Gln Ala Val Ala Asp Arg Leu Gln Arg Gln
100 105 110

Glu Glu Glu Arg Met Asn Cys Ser Pro Thr Ser Gln Ile Asp Asn Ile
115 120 125

Gly Glu Glu Glu Met Asp Ala Ser Thr Thr His His Lys Arg Lys Thr
130 135 140

Met Asn Asp Phe Asp Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe Gly
145 150 155 160

Lys Val Ile Leu Val Arg Glu Lys Ala Ser Gly Lys Tyr Tyr Ala Met
165 170 175

Lys Ile Leu Lys Lys Glu Val Ile Ile Ala Lys Val Thr Asp Leu Leu
180 185 190

Lys Leu Ile Thr Lys Phe Leu Phe Ala Val Cys Met Cys Leu Trp Ala
195 200 205

His Glu Phe Thr Cys
210

<210> 157

<211> 352

<212> PRT

<213> Homo sapiens

<400> 157

Met Gly Gly Lys Pro Ala Asn Arg Met Met Pro Tyr Pro Phe Pro Ser
1 5 10 15

Gly Thr Trp Lys Val Lys Trp Val Ala Ser Arg Asn Ala Phe Lys Pro
20 25 30

Arg	Ile	Gly	Ile	Leu	Ile	Lys	Thr	Leu	Ile	Tyr	Ser	Ser	Gln	Phe	Pro	35	40	45
Leu	Gly	Asn	Leu	Glu	Lys	Ile	Ser	Gln	Leu	Leu	Ser	Lys	Ser	Ala	Gln	50	55	60
Cys	Pro	Leu	Arg	Val	His	Tyr	Leu	Ser	Ser	Gln	Tyr	Gly	Asp	Glu	Arg	65	70	75
Cys	Phe	Met	Phe	Val	Leu	Ile	Ser	Pro	Thr	Lys	Ser	Val	Ile	Ile	Thr	85	90	95
Ile	Leu	Ser	Leu	Leu	Phe	Thr	Leu	Gln	Leu	Phe	Phe	His	Leu	Ser	Arg	100	105	110
Glu	Arg	Val	Phe	Ser	Glu	Asp	Arg	Thr	Arg	Phe	Tyr	Gly	Ala	Glu	Ile	115	120	125
Val	Ser	Ala	Leu	Asp	Tyr	Leu	His	Ser	Gly	Lys	Ile	Val	Tyr	Arg	Asp	130	135	140
Leu	Lys	Leu	Glu	Asn	Leu	Met	Leu	Asp	Lys	Asp	Gly	His	Ile	Lys	Ile	145	150	155
Thr	Asp	Phe	Gly	Leu	Cys	Lys	Glu	Gly	Ile	Thr	Asp	Ala	Ala	Thr	Met	165	170	175
Lys	Thr	Phe	Cys	Gly	Thr	Pro	Glu	Tyr	Leu	Ala	Pro	Glu	Val	Leu	Glu	180	185	190
Asp	Asn	Asp	Tyr	Gly	Arg	Ala	Val	Asp	Trp	Trp	Gly	Leu	Gly	Val	Val	195	200	205
Met	Tyr	Glu	Met	Met	Cys	Gly	Arg	Leu	Pro	Phe	Tyr	Asn	Gln	Asp	His	210	215	220
Glu	Lys	Leu	Phe	Glu	Leu	Ile	Leu	Met	Glu	Asp	Ile	Lys	Phe	Pro	Arg	225	230	235
Thr	Leu	Ser	Ser	Asp	Ala	Lys	Ser	Leu	Leu	Ser	Gly	Leu	Leu	Ile	Lys	245	250	255
Asp	Pro	Asn	Lys	Arg	Leu	Gly	Gly	Gly	Pro	Asp	Asp	Ala	Lys	Glu	Ile	260	265	270
Met	Arg	His	Ser	Phe	Phe	Ser	Gly	Val	Asn	Trp	Gln	Asp	Val	Tyr	Asp	275	280	285
Lys	Lys	Leu	Val	Pro	Pro	Phe	Lys	Pro	Gln	Val	Thr	Ser	Glu	Thr	Asp	290	295	300
Thr	Arg	Tyr	Phe	Asp	Glu	Glu	Phe	Thr	Ala	Gln	Thr	Ile	Thr	Ile	Thr	305	310	315
Pro	Pro	Glu	Lys	Tyr	Asp	Glu	Asp	Gly	Met	Asp	Cys	Met	Asp	Asn	Glu	325	330	335

Arg Arg Pro His Phe Pro Gln Phe Ser Tyr Ser Ala Ser Gly Arg Glu
 340 345 350

<210> 158
 <211> 132
 <212> PRT
 <213> Homo sapiens

<400> 158

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
 1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Gly Asn Ser Lys Lys
 20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
 35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His His Ser His
 50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys
 65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
 85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
 100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Val Cys
 115 120 125

Met Thr Leu Glu
 130

<210> 159
 <211> 192
 <212> PRT
 <213> Homo sapiens

<400> 159

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
 1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Gly Asn Ser Lys Lys
 20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
 35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His His Ser His
 50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys

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65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Ile Asp
115 120 125

Lys Glu Ile Glu Leu His Arg Ile Leu His His Lys His Val Val Gln
130 135 140

Phe Tyr His Tyr Phe Glu Asp Lys Glu Asn Ile Tyr Ile Leu Leu Glu
145 150 155 160

Tyr Cys Ser Arg Arg Val Ser Val Asn Ser Tyr Leu Arg Thr Phe Ala
165 170 175

Tyr Pro Glu Leu Thr Trp Tyr Ser Lys Ser Ile Leu Ser Gly Ile Thr
180 185 190

<210> 160
<211> 207
<212> PRT
<213> Homo sapiens

<400> 160

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Gly Asn Ser Lys Lys
20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His His Ser His
50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys
65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Ile Asp
115 120 125

Lys Glu Ile Glu Leu His Arg Ile Leu His His Lys His Val Val Gln
130 135 140

1000
900
800
700
600
500
400
300
200
100
0

Gly Pro Gly Ser Arg Pro Thr Ala Val Glu Gly Leu Ala Leu Gly Ser
 210 215 220
 Ser Ala Ser Ser Ser Ser Glu Pro Pro Gln Ile Ile Ile Asn Pro Ala
 225 230 235 240
 Arg Gln Lys Met Val Gln Lys Leu Ala Leu Tyr Glu Asp Gly Ala Leu
 245 250 255
 Asp Ser Leu Gln Leu Leu Ser Ser Ser Ser Leu Pro Gly Leu Gly Leu
 260 265 270
 Glu Gln Asp Arg Gln Gly Pro Lys Lys Val Met Asn Phe Arg Ala Asp
 275 280 285
 Val Phe Thr Trp Ala Asp Pro Pro Asn Pro Glu Val Lys Val Leu Met
 290 295 300
 Val Arg Ser Ser His Gly Ala Arg Val Leu Ser Thr Leu Pro Ala Val
 305 310 315 320
 Gly Val Gly Ala His Ala Arg Trp Gly Glu Lys Glu Val Ala Leu Leu
 325 330 335

Phe

<210> 162
 <211> 122
 <212> PRT
 <213> Homo sapiens

<400> 162

Met Gly His Ala Leu Cys Val Cys Ser Arg Gly Thr Val Ile Ile Asp
 1 5 10 15
 Asn Lys Arg Tyr Leu Phe Ile Gln Lys Leu Gly Glu Gly Gly Phe Ser
 20 25 30
 Tyr Val Asp Leu Val Glu Gly Leu His Asp Gly His Phe Tyr Ala Leu
 35 40 45
 Lys Arg Ile Leu Cys His Glu Gln Gln Asp Arg Glu Glu Ala Gln Arg
 50 55 60
 Glu Ala Asp Met His Arg Leu Phe Asn His Pro Asn Ile Leu Arg Leu
 65 70 75 80
 Val Ala Tyr Cys Leu Arg Glu Arg Gly Ala Lys His Glu Ala Trp Leu
 85 90 95
 Leu Leu Pro Phe Phe Lys Val Arg Lys Thr Pro Val Tyr Gly Gly Gly
 100 105 110
 Cys Ser Arg Ala Thr Tyr Ser Arg Ala Val

115

120

<210> 163
 <211> 842
 <212> PRT
 <213> Homo sapiens

<400> 163

Met	Glu	Arg	Ala	Ile	Ser	Pro	Gly	Leu	Leu	Val	Arg	Ala	Leu	Leu	Leu
1				5					10					15	
Leu	Leu	Leu	Leu	Gly	Leu	Ala	Ala	Arg	Thr	Val	Ala	Ala	Gly	Arg	Ala
			20					25					30		
Arg	Gly	Leu	Pro	Ala	Pro	Thr	Ala	Glu	Ala	Ala	Phe	Gly	Leu	Gly	Ala
			35				40					45			
Ala	Ala	Ala	Pro	Thr	Ser	Ala	Thr	Arg	Val	Pro	Ala	Ala	Gly	Ala	Val
			50			55					60				
Ala	Ala	Ala	Glu	Val	Thr	Val	Glu	Asp	Ala	Glu	Ala	Leu	Pro	Ala	Ala
65					70				75					80	
Ala	Gly	Glu	Gln	Glu	Pro	Arg	Gly	Pro	Glu	Pro	Asp	Asp	Glu	Thr	Glu
			85					90						95	
Leu	Arg	Pro	Arg	Gly	Arg	Ser	Leu	Val	Ile	Ile	Ser	Thr	Leu	Asp	Gly
			100					105					110		
Arg	Ile	Ala	Ala	Leu	Asp	Pro	Glu	Asn	His	Gly	Lys	Lys	Gln	Trp	Asp
			115				120					125			
Leu	Asp	Val	Gly	Ser	Gly	Ser	Leu	Val	Ser	Ser	Ser	Leu	Ser	Lys	Pro
			130			135					140				
Glu	Val	Phe	Gly	Asn	Lys	Met	Ile	Ile	Pro	Ser	Leu	Asp	Gly	Ala	Leu
145					150					155				160	
Phe	Gln	Trp	Asp	Arg	Asp	Arg	Glu	Ser	Met	Glu	Thr	Val	Pro	Phe	Thr
			165					170						175	
Val	Glu	Ser	Leu	Leu	Glu	Ser	Ser	Tyr	Lys	Phe	Gly	Asp	Asp	Val	Val
			180					185					190		
Leu	Val	Gly	Gly	Lys	Ser	Leu	Thr	Thr	Tyr	Gly	Leu	Ser	Ala	Tyr	Ser
			195				200					205			
Gly	Lys	Val	Arg	Tyr	Ile	Cys	Ser	Ala	Leu	Gly	Cys	Arg	Gln	Trp	Asp
			210			215					220				
Ser	Asp	Glu	Met	Glu	Gln	Glu	Glu	Asp	Ile	Leu	Leu	Leu	Gln	Arg	Thr
225					230					235				240	
Gln	Lys	Thr	Val	Arg	Ala	Val	Gly	Pro	Arg	Ser	Gly	Asn	Glu	Lys	Trp
				245					250					255	

Asn	Phe	Ser	Val	Gly	His	Phe	Glu	Leu	Arg	Tyr	Ile	Pro	Asp	Met	Glu	260	265	270
Thr	Arg	Ala	Gly	Phe	Ile	Glu	Ser	Thr	Phe	Lys	Pro	Asn	Glu	Asn	Thr	275	280	285
Glu	Glu	Ser	Lys	Ile	Ile	Ser	Asp	Val	Glu	Glu	Gln	Glu	Ala	Ala	Ile	290	295	300
Met	Asp	Ile	Val	Ile	Lys	Val	Ser	Val	Ala	Asp	Trp	Lys	Val	Met	Ala	305	310	315
Phe	Ser	Lys	Lys	Gly	Gly	His	Leu	Glu	Trp	Glu	Tyr	Gln	Phe	Cys	Thr	325	330	335
Pro	Ile	Ala	Ser	Ala	Trp	Leu	Leu	Lys	Asp	Gly	Lys	Val	Ile	Pro	Ile	340	345	350
Ser	Leu	Phe	Asp	Asp	Thr	Ser	Tyr	Thr	Ser	Asn	Asp	Asp	Val	Leu	Glu	355	360	365
Asp	Glu	Glu	Asp	Ile	Val	Glu	Ala	Ala	Arg	Gly	Ala	Thr	Glu	Asn	Ser	370	375	380
Val	Tyr	Leu	Gly	Met	Tyr	Arg	Gly	Gln	Leu	Tyr	Leu	Gln	Ser	Ser	Val	385	390	395
Arg	Ile	Ser	Glu	Lys	Phe	Pro	Ser	Ser	Pro	Lys	Ala	Leu	Glu	Ser	Val	405	410	415
Thr	Asn	Glu	Asn	Ala	Ile	Ile	Pro	Leu	Pro	Thr	Ile	Lys	Trp	Lys	Pro	420	425	430
Leu	Ile	His	Ser	Pro	Ser	Arg	Thr	Pro	Val	Leu	Val	Gly	Ser	Asp	Glu	435	440	445
Phe	Asp	Lys	Cys	Leu	Ser	Asn	Asp	Lys	Phe	Ser	His	Glu	Glu	Tyr	Ser	450	455	460
Asn	Gly	Ala	Leu	Ser	Ile	Leu	Gln	Tyr	Pro	Tyr	Asp	Asn	Gly	Tyr	Tyr	465	470	475
Leu	Pro	Tyr	Tyr	Lys	Arg	Glu	Arg	Asn	Lys	Arg	Ser	Thr	Gln	Ile	Thr	485	490	495
Val	Arg	Phe	Leu	Asp	Asn	Pro	His	Tyr	Asn	Lys	Asn	Ile	Arg	Lys	Lys	500	505	510
Asp	Pro	Val	Leu	Leu	Leu	His	Trp	Trp	Lys	Glu	Ile	Val	Ala	Thr	Ile	515	520	525
Leu	Phe	Cys	Ile	Ile	Ala	Thr	Thr	Phe	Ile	Val	Arg	Arg	Leu	Phe	His	530	535	540
Pro	His	Pro	His	Arg	Gln	Arg	Lys	Glu	Ser	Glu	Thr	Gln	Cys	Gln	Thr	545	550	555

Glu	Asn	Lys	Tyr	Asp	Ser	Val	Ser	Gly	Glu	Ala	Asn	Asp	Ser	Ser	Trp	
				565					570					575		
Asn	Asp	Ile	Lys	Asn	Ser	Gly	Tyr	Ile	Ser	Arg	Tyr	Leu	Thr	Asp	Phe	
				580					585					590		
Glu	Pro	Ile	Gln	Cys	Leu	Gly	Arg	Gly	Gly	Phe	Gly	Val	Val	Phe	Glu	
				595					600					605		
Ala	Lys	Asn	Lys	Val	Asp	Asp	Cys	Asn	Tyr	Ala	Ile	Lys	Arg	Ile	Arg	
				610					615					620		
Leu	Pro	Asn	Arg	Glu	Leu	Ala	Arg	Glu	Lys	Val	Met	Arg	Glu	Val	Lys	
				625					630					635		
Ala	Leu	Ala	Lys	Leu	Glu	His	Pro	Gly	Ile	Val	Arg	Tyr	Phe	Asn	Ala	
				645					650					655		
Trp	Leu	Glu	Ala	Pro	Pro	Glu	Lys	Trp	Gln	Glu	Lys	Met	Asp	Glu	Ile	
				660					665					670		
Trp	Leu	Lys	Asp	Glu	Ser	Thr	Asp	Trp	Pro	Leu	Ser	Ser	Pro	Ser	Pro	
				675					680					685		
Met	Asp	Ala	Pro	Ser	Val	Lys	Ile	Arg	Arg	Met	Asp	Pro	Phe	Ser	Thr	
				690					695					700		
Lys	Glu	His	Ile	Glu	Ile	Ile	Ala	Pro	Ser	Pro	Gln	Arg	Ser	Arg	Ser	
				705					710					715		
Phe	Ser	Val	Gly	Ile	Ser	Cys	Asp	Gln	Thr	Ser	Ser	Ser	Glu	Ser	Gln	
				725					730					735		
Phe	Ser	Pro	Leu	Glu	Phe	Ser	Gly	Met	Asp	His	Glu	Asp	Ile	Ser	Glu	
				740					745					750		
Ser	Val	Asp	Ala	Ala	Tyr	Asn	Leu	Gln	Asp	Ser	Cys	Leu	Thr	Asp	Cys	
				755					760					765		
Asp	Val	Glu	Asp	Gly	Thr	Met	Asp	Gly	Asn	Asp	Glu	Gly	His	Ser	Phe	
				770					775					780		
Glu	Leu	Cys	Pro	Ser	Glu	Ala	Ser	Pro	Tyr	Val	Arg	Ser	Arg	Glu	Arg	
				785					790					795		
Thr	Ser	Ser	Ser	Ile	Val	Phe	Glu	Asp	Ser	Gly	Cys	Asp	Asn	Ala	Ser	
				805					810					815		
Ser	Lys	Glu	Glu	Pro	Lys	Thr	Asn	Arg	Leu	His	Ile	Gly	Asn	His	Cys	
				820					825					830		
Ala	Asn	Lys	Leu	Thr	Val	Thr	Val	Leu	Phe							
				835					840							

<210> 164

<211> 743

<212> PRT

Met 1	Gly	Ser	Arg	Ala 5	Gln	Lys	Ser	Ala	Gly 10	Asn	Ala	Glu	Leu	Trp 15	Glu
Pro	Leu	Pro	Glu 20	Gly	Arg	Pro	Arg	Pro	Ala 25	Gly	Thr	Ser	Ser 30	Ala	Val
Ser	Ala	Trp 35	Ala	Ser	Leu	Lys	Leu 40	Cys	Leu	Arg	Gly	Gly 45	Ser	Gly	Arg
Arg 50	Gln	Arg	Leu	Gly	Gly	Gly 55	Arg	Met	Gln	Pro	Glu 60	Glu	Gly	His	Arg
Leu 65	Ala	Ala	Gly	Ala	Ala 70	Val	Arg	Gly	Ala 75	Ala	Ala	Thr	Val	Leu	Leu 80
Arg	Leu	Arg	Asp 85	Asp	Leu	Asn	Val	Thr 90	Arg	Leu	Ser	His	Phe 95	Glu	Tyr
Val	Lys	Asn 100	Glu	Asp	Leu	Glu	Lys 105	Ile	Gly	Met	Gly	Arg	Pro 110	Gly	Gln
Arg	Arg	Leu 115	Trp	Glu	Ala	Val	Lys 120	Arg	Arg	Lys	Ala	Leu 125	Cys	Lys	Arg
Lys 130	Ser	Trp	Met	Asn	Lys	Val 135	Phe	Ser	Gly	Lys	Arg 140	Leu	Glu	Ala	Glu
Phe 145	Pro	Pro	His	His	Ser 150	Gln	Ser	Thr	Phe	Arg 155	Lys	Thr	Ser	Pro	Ala 160
Pro	Gly	Gly	Pro 165	Ala	Gly	Glu	Gly	Pro	Leu 170	Gln	Ser	Leu	Thr	Cys 175	Leu
Ile	Gly	Glu 180	Lys	Asp	Leu	Arg	Leu 185	Leu	Glu	Lys	Leu	Gly 190	Asp	Gly	Ser
Phe 195	Gly	Val	Val	Arg	Arg	Gly	Glu 200	Trp	Asp	Ala	Pro 205	Ser	Gly	Lys	Thr
Val 210	Ser	Pro	Pro	Gln	Pro	Ala 215	Phe	Phe	Thr	Gln	Lys 220	Pro	Thr	Tyr	Asp
Pro 225	Val	Ser	Glu	Asp	Gln 230	Asp	Pro	Leu	Ser	Ser 235	Asp	Phe	Lys	Arg	Leu 240
Gly	Leu	Arg	Lys 245	Pro	Gly	Leu	Pro	Arg	Gly 250	Leu	Trp	Leu	Ala	Lys 255	Pro
Ser	Ala	Arg 260	Val	Pro	Gly	Thr	Lys 265	Ala	Ser	Arg	Gly	Ser	Gly 270	Ala	Glu
Val 275	Thr	Leu	Ile	Asp	Phe	Gly 280	Glu	Glu	Pro	Val	Val 285	Pro	Ala	Leu	Arg

Pro Cys Ala Pro Ser Leu Ala Gln Leu Ala Met Asp Ala Cys Ser Leu
290 295 300

Leu Asp Glu Thr Pro Pro Gln Ser Pro Thr Arg Ala Leu Pro Arg Pro
305 310 315 320

Leu His Pro Thr Pro Val Val Asp Trp Asp Ala Arg Pro Leu Pro Pro
325 330 335

Pro Pro Ala Tyr Asp Asp Val Ala Gln Asp Glu Asp Asp Phe Glu Ile
340 345 350

Cys Ser Ile Asn Ser Thr Leu Val Gly Ala Gly Val Pro Ala Gly Pro
355 360 365

Ser Gln Gly Gln Thr Asn Tyr Ala Phe Val Pro Glu Gln Ala Arg Pro
370 375 380

Pro	Pro	Pro	Leu	Glu	Asp	Asn	Leu	Phe	Leu	Pro	Pro	Gln	Gly	Gly	Gly
385					390					395					400

Lys Pro Pro Ser Ser Ala Gln Thr Ala Glu Ile Phe Gln Ala Leu Gln
405 410 415

Gln Glu Cys Met Arg Gln Leu Gln Ala Pro Ala Gly Ser Pro Ala Pro
 420 425 430

Ser	Pro	Ser	Pro	Gly	Gly	Asp	Asp	Lys	Pro	Gln	Val	Pro	Pro	Arg	Val
		435					440					445			

Pro Ile Pro Pro Arg Pro Thr Arg Pro His Val Gln Leu Ser Pro Ala
450 455 460

Pro Pro Gly Glu Glu Glu Thr Ser Gln Trp Pro Gly Pro Ala Ser Pro
465 470 475 480

Pro Arg Val Pro Pro Arg Glu Pro Leu Ser Pro Gln Gly Ser Arg Thr
485 490 495

Pro Ser Pro Leu Val Pro Pro Gly Ser Ser Pro Leu Pro Pro Arg Leu
500 505 510

Ser Ser Ser Pro Gly Lys Thr Met Pro Thr Thr Gln Ser Phe Ala Ser
515 520 525

Asp Pro Lys Tyr Ala Thr Pro Gln Val Ile Gln Ala Pro Gly Pro Arg
530 535 540

Ala Gly Pro Cys Ile Leu Pro Ile Val Arg Asp Gly Lys Lys Val Ser
545 550 555 560

Ser Thr His Tyr Tyr Leu Leu Pro Glu Arg Pro Ser Tyr Leu Glu Arg
565 570 575

Tyr Gln Arg Phe Leu Arg Glu Ala Gln Ser Pro Glu Glu Pro Thr Pro
580 585 590

Leu Pro Val Pro Leu Leu Leu Pro Pro Pro Ser Thr Pro Ala Pro Ala
595 600 605

Ala Pro Thr Ala Thr Val Arg Pro Met Pro Gln Ala Ala Leu Asp Pro
610 615 620

Lys Ala Asn Phe Ser Thr Asn Asn Ser Asn Pro Gly Ala Arg Pro Pro
625 630 635 640

Pro Pro Arg Ala Thr Ala Arg Leu Pro Gln Arg Gly Cys Pro Gly Asp
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Gly Pro Glu Ala Gly Arg Pro Ala Asp Lys Ile Gln Met Ala Met Val
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His Gly Val Thr Thr Glu Glu Cys Gln Ala Ala Leu Gln Cys His Gly
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Trp Ser Val Gln Arg Ala Cys Pro Val Ser Glu Gly Gly Ala Ala Leu
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Arg Ala Gly Ser Ala Ala Gln Arg Glu Cys His Lys Val Leu Glu Met
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Phe Asp Trp Asn Leu Glu Gln Ala Gly Cys His Leu Leu Gly Ser Trp
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Gly Pro Ala His His Lys Arg
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Gly Asp Ser Arg Ser Leu Pro Phe Ser Glu Asn Val Ser Ala Val Gln
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Lys Leu Asp Phe Ser Asp Thr Met Val Gln Gln Lys Leu Asp Asp Ile
35 40 45

Lys Asp Arg Ile Lys Arg Glu Ile Arg Lys Glu Leu Lys Ile Lys Glu
50 55 60

Gly Ala Glu Asn Leu Arg Lys Val Thr Thr Asp Lys Lys Ser Leu Ala
65 70 75 80

Tyr Val Asp Asn Ile Leu Lys Lys Ser Asn Lys Lys Leu Glu Glu Leu
85 90 95

His His Lys Leu Gln Glu Leu Asn Ala His Ile Val Val Ser Asp Pro

165
604
PRT
Homo sapiens
165

100					105					110						
Glu	Asp	Ile	Thr	Asp	Cys	Pro	Arg	Thr	Pro	Asp	Thr	Pro	Asn	Asn	Asp	
115					120					125						
Pro	Arg	Cys	Ser	Thr	Ser	Asn	Asn	Arg	Leu	Lys	Ala	Leu	Gln	Lys	Gln	
130					135					140						
Leu	Asp	Ile	Glu	Leu	Lys	Val	Lys	Gln	Gly	Ala	Glu	Asn	Met	Ile	Gln	
145					150					155					160	
Met	Tyr	Ser	Asn	Gly	Ser	Ser	Lys	Asp	Arg	Lys	Leu	His	Gly	Thr	Ala	
165					170					175						
Gln	Gln	Leu	Leu	Gln	Asp	Ser	Lys	Thr	Lys	Ile	Glu	Val	Ile	Arg	Met	
180					185					190						
Gln	Ile	Leu	Gln	Ala	Val	Gln	Thr	Asn	Glu	Leu	Ala	Phe	Asp	Asn	Ala	
195					200					205						
Lys	Pro	Val	Ile	Ser	Pro	Leu	Glu	Leu	Arg	Met	Glu	Glu	Leu	Arg	His	
210					215					220						
His	Phe	Arg	Ile	Glu	Phe	Ala	Val	Ala	Glu	Gly	Ala	Lys	Asn	Val	Met	
225					230					235					240	
Lys	Leu	Leu	Gly	Ser	Gly	Lys	Val	Thr	Asp	Arg	Lys	Ala	Leu	Ser	Glu	
245					250					255						
Ala	Gln	Ala	Arg	Phe	Asn	Glu	Ser	Ser	Gln	Lys	Leu	Asp	Leu	Leu	Lys	
260					265					270						
Tyr	Ser	Leu	Glu	Gln	Arg	Leu	Asn	Glu	Val	Pro	Lys	Asn	His	Pro	Lys	
275					280					285						
Ser	Arg	Ile	Ile	Ile	Glu	Glu	Leu	Ser	Leu	Val	Ala	Ala	Ser	Pro	Thr	
290					295					300						
Leu	Ser	Pro	Arg	Gln	Ser	Met	Ile	Ser	Thr	Gln	Asn	Gln	Tyr	Ser	Thr	
305					310					315					320	
Leu	Ser	Lys	Pro	Ala	Ala	Leu	Thr	Gly	Thr	Leu	Glu	Val	Arg	Leu	Met	
325					330					335						
Gly	Cys	Gln	Asp	Ile	Leu	Glu	Asn	Val	Pro	Gly	Arg	Ser	Lys	Ala	Thr	
340					345					350						
Ser	Val	Ala	Leu	Pro	Gly	Trp	Ser	Pro	Ser	Glu	Thr	Arg	Ser	Ser	Phe	
355					360					365						
Met	Ser	Arg	Thr	Ser	Lys	Ser	Lys	Ser	Gly	Ser	Ser	Arg	Asn	Leu	Leu	
370					375					380						
Lys	Thr	Asp	Asp	Leu	Ser	Asn	Asp	Val	Cys	Ala	Val	Leu	Lys	Leu	Asp	
385					390					395					400	
Asn	Thr	Val	Val	Gly	Gln	Thr	Ser	Trp	Lys	Pro	Ile	Ser	Asn	Gln	Ser	

100 105 110
 115 120 125
 130 135 140
 145 150 155 160
 165 170 175
 180 185 190
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 210 215 220
 225 230 235 240
 245 250 255
 260 265 270
 275 280 285
 290 295 300
 305 310 315 320
 325 330 335
 340 345 350
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 370 375 380
 385 390 395 400

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	405	410	415
Trp Asp Gln Lys Phe Thr Leu Glu Leu Asp Arg Ser Arg Glu Leu Glu	420	425	430
Ile Ser Val Tyr Trp Arg Asp Trp Arg Ser Leu Cys Ala Val Lys Phe	435	440	445
Leu Arg Leu Glu Asp Phe Leu Asp Asn Gln Arg His Gly Met Cys Leu	450	455	460
Tyr Leu Glu Pro Gln Gly Thr Leu Phe Ala Glu Val Thr Phe Phe Asn	465	470	475
Pro Val Ile Glu Arg Arg Pro Lys Leu Gln Arg Gln Lys Lys Ile Phe	485	490	495
Ser Lys Gln Gln Gly Lys Thr Phe Leu Arg Ala Pro Gln Met Asn Ile	500	505	510
Asn Ile Ala Thr Trp Gly Arg Leu Val Arg Arg Ala Ile Pro Thr Val	515	520	525
Asn His Ser Gly Thr Phe Ser Pro Gln Ala Pro Val Pro Thr Thr Val	530	535	540
Pro Val Val Asp Val Arg Ile Pro Gln Leu Ala Pro Pro Ala Arg Tyr	545	550	555
Val Ser Glu Ile Leu Ser Ile Ser Tyr Thr Lys Leu Leu Gly His Ser	565	570	575
Tyr Val Leu Ile Ile Ala Gly Val Leu Ser Leu Ala Phe Phe Pro Ser	580	585	590
Ser Ile Leu Lys Val Val Phe Cys Leu Leu Lys Lys	595	600	
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Met Ala Ser Asn Pro Glu Arg Gly Glu Ile Leu Leu Thr Glu Leu Gln	1	5	10
Gly Asp Ser Arg Ser Leu Pro Phe Ser Glu Asn Val Ser Ala Val Gln	20	25	30
Lys Leu Asp Phe Ser Asp Thr Met Val Gln Gln Lys Leu Asp Asp Ile	35	40	45
Lys Asp Arg Ile Lys Arg Glu Ile Arg Lys Glu Leu Lys Ile Lys Glu	50	55	60

Gly	Ala	Glu	Asn	Leu	Arg	Lys	Val	Thr	Thr	Asp	Lys	Lys	Ser	Leu	Ala	
65					70					75					80	
Tyr	Val	Asp	Asn	Ile	Leu	Lys	Lys	Ser	Asn	Lys	Lys	Leu	Glu	Glu	Leu	
			85						90					95		
His	His	Lys	Leu	Gln	Glu	Leu	Asn	Ala	His	Ile	Val	Val	Ser	Asp	Pro	
			100					105						110		
Glu	Asp	Ile	Thr	Asp	Cys	Pro	Arg	Thr	Pro	Asp	Thr	Pro	Asn	Asn	Asp	
		115					120						125			
Pro	Arg	Cys	Ser	Thr	Ser	Asn	Asn	Arg	Leu	Lys	Ala	Leu	Gln	Lys	Gln	
	130					135					140					
Leu	Asp	Ile	Glu	Leu	Lys	Val	Lys	Gln	Gly	Ala	Glu	Asn	Met	Ile	Gln	
145					150					155					160	
Met	Tyr	Ser	Asn	Gly	Ser	Ser	Lys	Asp	Arg	Lys	Leu	His	Gly	Thr	Ala	
				165					170						175	
Gln	Gln	Leu	Leu	Gln	Asp	Ser	Lys	Thr	Lys	Ile	Glu	Val	Ile	Arg	Met	
			180					185						190		
Gln	Ile	Leu	Gln	Ala	Val	Gln	Thr	Asn	Glu	Leu	Ala	Phe	Asp	Asn	Ala	
		195					200						205			
Lys	Pro	Val	Ile	Ser	Pro	Leu	Glu	Leu	Arg	Met	Glu	Glu	Leu	Arg	His	
	210					215						220				
His	Phe	Arg	Ile	Glu	Phe	Ala	Val	Ala	Glu	Gly	Ala	Lys	Asn	Val	Met	
225					230					235					240	
Lys	Leu	Leu	Gly	Ser	Gly	Lys	Val	Thr	Asp	Arg	Lys	Ala	Leu	Ser	Glu	
			245						250					255		
Ala	Gln	Ala	Arg	Phe	Asn	Glu	Ser	Ser	Gln	Lys	Leu	Asp	Leu	Leu	Lys	
			260					265					270			
Tyr	Ser	Leu	Glu	Gln	Arg	Leu	Asn	Glu	Val	Pro	Lys	Asn	His	Pro	Lys	
		275					280						285			
Ser	Arg	Ile	Ile	Ile	Glu	Glu	Leu	Ser	Leu	Val	Ala	Ala	Ser	Pro	Thr	
		290				295					300					
Leu	Ser	Pro	Arg	Gln	Ser	Met	Ile	Ser	Thr	Gln	Asn	Gln	Tyr	Ser	Thr	
305					310					315					320	
Leu	Ser	Lys	Pro	Ala	Ala	Leu	Thr	Gly	Thr	Leu	Glu	Val	Arg	Leu	Met	
			325						330					335		
Gly	Cys	Gln	Asp	Ile	Leu	Glu	Asn	Val	Pro	Gly	Arg	Ser	Lys	Ala	Thr	
			340					345					350			
Ser	Val	Ala	Leu	Pro	Gly	Trp	Ser	Pro	Ser	Glu	Thr	Arg	Ser	Ser	Phe	
		355					360					365				

Met Ser Arg Thr Ser Lys Ser Lys Ser Gly Ser Ser Arg Asn Leu Leu
370 375 380

Lys Thr Asp Asp Leu Ser Asn Asp Val Cys Ala Val Leu Lys Leu Asp
385 390 395 400

Asn Thr Val Val Gly Gln Thr Ser Trp Lys Pro Ile Ser Asn Gln Ser
405 410 415

Trp Asp Gln Lys Phe Thr Leu Glu Leu Asp Arg Ser Arg Glu Leu Glu
420 425 430

Ile Ser Val Tyr Trp Arg Asp Trp Arg Ser Leu Cys Ala Val Lys Phe
435 440 445

Leu Arg Leu Glu Asp Phe Leu Asp Asn Gln Arg His Gly Met Cys Leu
450 455 460

Tyr Leu Glu Pro Gln Gly Thr Leu Phe Ala Glu Val Thr Phe Phe Asn
465 470 475 480

Pro Val Ile Glu Arg Arg Pro Lys Leu Gln Arg Gln Lys Lys Ile Phe
485 490 495

Ser Lys Gln Gln Gly Lys Thr Phe Leu Arg Ala Pro Gln Met Asn Ile
500 505 510

Asn Ile Ala Thr Trp Gly Arg Leu Val Arg Arg Ala Ile Pro Thr Val
515 520 525

Asn His Ser Gly Thr Phe Ser Pro Gln Ala Pro Val Pro Thr Thr Val
530 535 540

Pro Val Val Asp Val Arg Ile Pro Gln Leu Ala Pro Pro Ala Ser Asp
545 550 555 560

Ser Thr Val Thr Lys Leu Asp Phe Asp Leu Glu Pro Glu Pro Pro Pro
565 570 575

Ala Pro Pro Arg Ala Ser Ser Leu Gly Glu Ile Asp Glu Ser Ser Glu
580 585 590

Leu Arg Val Leu Asp Ile Pro Gly Gln Ala Ser His Phe Lys Pro Cys
595 600 605

Ile Ile Pro Leu His
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<210> 167

<211> 133

<212> PRT

<213> Homo sapiens

<400> 167

Met Val Ser Ser Gln Lys Leu Glu Lys Pro Ile Glu Met Gly Ser Ser
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Glu Pro Leu Pro Ile Ala Asp Gly Asp Arg Arg Arg Lys Lys Lys Arg
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Arg Gly Arg Ala Thr Asp Ser Leu Pro Gly Lys Phe Glu Asp Met Tyr
35 40 45

Lys Leu Thr Ser Glu Leu Leu Gly Glu Gly Ala Tyr Ala Lys Val Gln
50 55 60

Gly Ala Val Ser Leu Gln Asn Gly Lys Glu Tyr Ala Val Lys Ile Ile
65 70 75 80

Glu Lys Gln Ala Gly His Ser Arg Ser Arg Val Phe Arg Glu Val Glu
85 90 95

Thr Leu Tyr Gln Cys Gln Gly Asn Lys Asn Ile Leu Glu Leu Ile Glu
100 105 110

Phe Phe Glu Asp Asp Thr Arg Phe Tyr Leu Val Phe Glu Lys Leu Gln
115 120 125

Gly Gly Thr Tyr Arg
130

<210> 168

<211> 153

<212> PRT

<213> Homo sapiens

<400> 168

Met Leu Gln Val Gly Val Leu Arg Asp Arg Ser Pro Ala Gly Ala Ser
1 5 10 15

Glu Gly Phe His Val Arg Gly Arg Trp Arg Thr Glu Asp Cys His Leu
20 25 30

Arg Thr Lys Ala Ile Glu Thr Leu Arg Val Ala Gly Arg His Gln Leu
35 40 45

Pro Asp Arg Ser Phe Ile Ser Phe Gly Ile Ser Ser Leu Gln Met Val
50 55 60

Ser Ser Gln Lys Leu Glu Lys Pro Ile Glu Met Gly Ser Ser Glu Pro
65 70 75 80

Leu Pro Ile Ala Asp Gly Asp Arg Arg Arg Lys Lys Lys Arg Arg Gly
85 90 95

Arg Ala Thr Asp Ser Leu Pro Gly Lys Phe Glu Asp Met Tyr Lys Leu
100 105 110

Thr Ser Glu Leu Leu Gly Glu Gly Ala Tyr Ala Lys Val Gln Gly Ala
115 120 125

Val Ser Leu Gln Asn Gly Lys Glu Tyr Ala Val Lys Val Ser Val Ser

168 153 PRT Homo sapiens 168

130	135	140
Ala Glu Cys Gln Ala Leu Leu Cys Lys		
145	150	
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<211> 231		
<212> PRT		
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<400> 169		
Met Gly Ser Gly Met Lys Leu Asn Asn Ser Cys Thr Pro Ile Thr Thr		
1	5	10 15
Pro Glu Leu Thr Thr Pro Cys Gly Ser Ala Glu Tyr Met Ala Pro Glu		
	20	25 30
Val Val Glu Val Phe Thr Asp Gln Ala Thr Phe Tyr Asp Lys Arg Cys		
	35	40 45
Asp Leu Trp Ser Leu Gly Val Val Leu Tyr Ile Met Leu Ser Gly Tyr		
	50	55 60
Pro Pro Phe Val Gly His Cys Gly Ala Asp Cys Gly Trp Asp Arg Gly		
65	70	75 80
Glu Val Cys Arg Val Cys Gln Asn Lys Leu Phe Glu Ser Ile Gln Glu		
	85	90 95
Gly Lys Tyr Glu Phe Pro Asp Lys Asp Trp Ala His Ile Ser Ser Glu		
	100	105 110
Ala Lys Asp Leu Ile Ser Lys Leu Leu Val Arg Asp Ala Lys Gln Arg		
	115	120 125
Leu Ser Ala Ala Gln Val Leu Gln His Pro Trp Val Gln Gly Gln Ala		
	130	135 140
Pro Glu Lys Gly Leu Pro Thr Pro Gln Val Leu Gln Arg Asn Ser Ser		
145	150	155 160
Thr Met Asp Leu Thr Leu Phe Ala Ala Glu Ala Ile Ala Leu Asn Arg		
	165	170 175
Gln Leu Ser Gln His Glu Glu Asn Glu Leu Ala Glu Glu Pro Glu Ala		
	180	185 190
Leu Ala Asp Gly Leu Cys Ser Met Lys Leu Ser Pro Pro Cys Lys Ser		
	195	200 205
Arg Leu Ala Arg Arg Arg Ala Leu Ala Gln Ala Gly Arg Gly Glu Asp		
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Arg Ser Pro Pro Thr Ala Leu		
225	230	

<210> 170
 <211> 146
 <212> PRT
 <213> Homo sapiens

<400> 170

Met Arg Lys Gly Val Leu Lys Asp Pro Glu Ile Ala Asp Leu Phe Tyr
 1 5 10 15

Lys Asp Asp Pro Glu Glu Leu Phe Ile Gly Leu His Glu Ile Gly His
 20 25 30

Gly Ser Phe Gly Ala Val Tyr Phe Ala Thr Asn Ala His Thr Ser Glu
 35 40 45

Val Val Ala Ile Lys Lys Met Ser Tyr Ser Gly Lys Gln Thr His Glu
 50 55 60

Lys Trp Gln Asp Ile Leu Lys Glu Val Lys Phe Leu Arg Gln Leu Lys
 65 70 75 80

His Pro Asn Thr Ile Glu Tyr Lys Gly Cys Tyr Leu Lys Glu His Thr
 85 90 95

Ala Trp Leu Val Met Glu Tyr Cys Leu Gly Ser Ala Ser Asp Leu Leu
 100 105 110

Glu Val His Lys Lys Pro Leu Gln Glu Val Glu Ile Ala Ala Ile Thr
 115 120 125

His Gly Ala Leu His Gly Leu Ala Tyr Leu His Ser His Ala Leu Ile
 130 135 140

His Arg
 145

<210> 171
 <211> 123
 <212> PRT
 <213> Homo sapiens

<400> 171

Met Met Glu Glu Leu His Ser Leu Asp Pro Arg Arg Gln Glu Leu Leu
 1 5 10 15

Glu Ala Arg Phe Thr Gly Val Gly Val Ser Lys Gly Pro Leu Asn Ser
 20 25 30

Glu Ser Ser Asn Gln Ser Leu Cys Ser Val Gly Ser Leu Ser Asp Lys
 35 40 45

Glu Val Glu Thr Pro Glu Lys Lys Gln Asn Asp Gln Arg Asn Arg Lys
 50 55 60

Arg Lys Ala Glu Pro Tyr Glu Thr Ser Gln Gly Lys Gly Thr Pro Arg

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65		70		75		80									
Gly	His	Lys	Ile	Ser	Asp	Tyr	Phe	Glu	Thr	Ala	Pro	Leu	Trp	Phe	Arg
				85					90					95	
Trp	Gln	Cys	Cys	Lys	Gly	Gly	Asn	Arg	Gly	Ala	Val	Cys	Ser	Ala	Asn
			100					105					110		
Pro	His	Val	Ser	Asp	Ala	Ser	Lys	Thr	Ser	Ala					
		115					120								
<210> 172															
<211> 478															
<212> PRT															
<213> Homo sapiens															
<400> 172															
Met	Val	Gly	Ile	Lys	Glu	Arg	Pro	Ser	Ser	Asn	Leu	Pro	Cys	Pro	Pro
1				5					10					15	
Leu	Pro	Pro	Gln	Thr	Gln	Ala	Cys	Pro	Pro	Leu	Ser	Trp	Pro	Gln	Arg
			20					25					30		
Leu	Asp	Ile	Leu	Leu	Gly	Thr	Ala	Arg	Ala	Ile	Gln	Phe	Leu	His	Gln
		35					40					45			
Asp	Ser	Pro	Ser	Leu	Ile	His	Gly	Asp	Ile	Lys	Ser	Ser	Asn	Val	Leu
	50					55					60				
Leu	Asp	Glu	Arg	Leu	Thr	Pro	Lys	Leu	Gly	Asp	Phe	Gly	Leu	Ala	Arg
65					70					75					80
Phe	Ser	Arg	Phe	Ala	Gly	Ser	Ser	Pro	Ser	Gln	Ser	Ser	Met	Val	Ala
				85					90					95	
Arg	Thr	Gln	Thr	Val	Arg	Gly	Thr	Leu	Ala	Tyr	Leu	Pro	Glu	Glu	Tyr
			100					105					110		
Ile	Lys	Thr	Gly	Arg	Leu	Ala	Val	Asp	Thr	Asp	Thr	Phe	Ser	Phe	Gly
		115					120					125			
Val	Val	Val	Leu	Glu	Thr	Leu	Ala	Gly	Gln	Arg	Ala	Val	Lys	Thr	His
		130					135					140			
Gly	Ala	Arg	Thr	Lys	Tyr	Leu	Lys	Asp	Leu	Val	Glu	Glu	Glu	Ala	Glu
145					150					155					160
Glu	Ala	Gly	Val	Ala	Leu	Arg	Ser	Thr	Gln	Ser	Thr	Leu	Gln	Ala	Gly
				165					170					175	
Leu	Ala	Ala	Asp	Ala	Trp	Ala	Ala	Pro	Ile	Ala	Met	Gln	Ile	Tyr	Lys
			180					185					190		
Lys	His	Leu	Asp	Pro	Arg	Pro	Gly	Pro	Cys	His	Leu	Ser	Trp	Ala	Trp
		195					200					205			

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Ala Trp Ala Ser Trp Pro Ala Ala Ala Cys Thr Ala Gly Pro Lys Gly
 210 215 220
 Arg Pro Pro Met Thr Gln Val Tyr Glu Arg Leu Glu Lys Leu Gln Ala
 225 230 235 240
 Val Val Ala Gly Val Pro Gly His Leu Glu Ala Ala Ser Cys Ile Pro
 245 250 255
 Phe Pro Gln Glu Asn Ser Tyr Val Ser Ser Thr Gly Arg Ala His Ser
 260 265 270
 Gly Ala Ala Pro Trp Gln Pro Leu Ala Ala Pro Ser Gly Ala Ser Ala
 275 280 285
 Gln Ala Ala Glu Gln Leu Gln Arg Gly Pro Asn Gln Pro Val Glu Ser
 290 295 300
 Asp Glu Ser Leu Gly Gly Leu Ser Ala Ala Leu Arg Ser Trp His Leu
 305 310 315 320
 Thr Pro Ser Cys Pro Leu Asp Pro Ala Pro Leu Arg Glu Ala Gly Cys
 325 330 335
 Pro Gln Gly Asp Thr Ala Gly Glu Ser Ser Trp Gly Ser Gly Pro Gly
 340 345 350
 Ser Arg Pro Thr Ala Val Glu Gly Leu Ala Leu Gly Ser Ser Ala Ser
 355 360 365
 Ser Ser Ser Glu Pro Pro Gln Ile Ile Ile Asn Pro Ala Arg Gln Lys
 370 375 380
 Met Val Gln Lys Leu Ala Leu Tyr Glu Asp Gly Ala Leu Asp Ser Leu
 385 390 395 400
 Gln Leu Leu Ser Ser Ser Ser Leu Pro Gly Leu Gly Leu Glu Gln Asp
 405 410 415
 Arg Gln Gly Pro Lys Lys Val Met Asn Phe Arg Ala Asp Val Phe Thr
 420 425 430
 Trp Ala Asp Pro Pro Asn Pro Glu Val Lys Val Leu Met Val Arg Ser
 435 440 445
 Ser His Gly Ala Arg Val Leu Ser Thr Leu Pro Ala Val Gly Val Gly
 450 455 460
 Ala His Ala Arg Trp Gly Glu Lys Glu Val Ala Leu Leu Phe
 465 470 475

<210> 173

<211> 344

<212> PRT

<213> Homo sapiens

<400> 173

Met	Ala	Gly	Gly	Pro	Gly	Pro	Gly	Glu	Pro	Ala	Ala	Pro	Gly	Ala	Gln	1	5	10	15
His	Phe	Leu	Tyr	Glu	Val	Pro	Pro	Trp	Val	Met	Cys	Arg	Phe	Tyr	Lys	20	25	30	
Val	Met	Asp	Ala	Leu	Glu	Pro	Ala	Asp	Trp	Cys	Gln	Phe	Ala	Ala	Leu	35	40	45	
Ile	Val	Arg	Asp	Gln	Thr	Glu	Leu	Arg	Leu	Cys	Glu	Arg	Ser	Gly	Gln	50	55	60	
Arg	Thr	Ala	Ser	Val	Leu	Trp	Pro	Trp	Ile	Asn	Arg	Asn	Ala	Arg	Val	65	70	75	80
Ala	Asp	Leu	Val	His	Ile	Leu	Thr	His	Leu	Gln	Leu	Leu	Arg	Ala	Arg	85	90	95	
Asp	Ile	Ile	Thr	Ala	Trp	His	Pro	Pro	Ala	Pro	Leu	Pro	Ser	Pro	Gly	100	105	110	
Thr	Thr	Ala	Pro	Arg	Pro	Ser	Ser	Ile	Pro	Ala	Pro	Ala	Glu	Ala	Glu	115	120	125	
Ala	Trp	Ser	Pro	Arg	Lys	Leu	Pro	Ser	Ser	Ala	Ser	Thr	Phe	Leu	Ser	130	135	140	
Pro	Ala	Phe	Pro	Gly	Ser	Gln	Thr	His	Ser	Gly	Pro	Glu	Leu	Gly	Leu	145	150	155	160
Val	Pro	Ser	Pro	Ala	Ser	Leu	Trp	Pro	Pro	Pro	Pro	Ser	Pro	Ala	Pro	165	170	175	
Ser	Ser	Thr	Lys	Pro	Gly	Pro	Glu	Ser	Ser	Val	Ser	Leu	Leu	Gln	Gly	180	185	190	
Ala	Arg	Pro	Ser	Pro	Phe	Cys	Trp	Pro	Leu	Cys	Glu	Ile	Ser	Arg	Gly	195	200	205	
Thr	His	Asn	Phe	Ser	Glu	Glu	Leu	Lys	Ile	Gly	Glu	Gly	Gly	Phe	Gly	210	215	220	
Cys	Val	Tyr	Arg	Ala	Val	Met	Arg	Asn	Thr	Val	Tyr	Ala	Val	Lys	Arg	225	230	235	240
Leu	Lys	Glu	Asn	Ala	Asp	Leu	Glu	Trp	Thr	Ala	Val	Lys	Gln	Ser	Phe	245	250	255	
Leu	Thr	Glu	Val	Glu	Gln	Leu	Ser	Arg	Phe	Arg	His	Pro	Asn	Ile	Val	260	265	270	
Asp	Phe	Ala	Gly	Tyr	Cys	Ala	Gln	Asn	Gly	Phe	Tyr	Cys	Leu	Val	Tyr	275	280	285	
Gly	Phe	Leu	Pro	Asn	Gly	Ser	Leu	Glu	Asp	Arg	Leu	His	Cys	Gln	Thr	290	295	300	

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Val Ala Ile Thr Ser Asp Asp Glu Ser Gln Ala Met Gln Thr Val Gly
 225 230 235 240

Val His Ser Ile Val Gln Gln Leu His Arg Asn Ser Ile Gln Phe Thr
 245 250 255

Asp Gly Tyr Glu Val Lys Glu Asp Ile Gly Val Gly Ser Tyr Ser Val
 260 265 270

Cys Lys Arg Cys Ile His Lys Ala Thr Asn Met Glu Phe Ala Val Lys
 275 280 285

Val Asn Phe Phe Tyr Leu Lys Cys Asn Ser Tyr Ser Ser Cys Ser Cys
 290 295 300

Met Ser Val Pro Val Lys Asn Tyr Thr Pro Leu Val Val Lys Ser Ala
 305 310 315 320

Phe Cys Tyr Lys Lys Val Lys Tyr Leu Ala Ser Asp Leu Gln Arg Ser
 325 330 335

<210> 175

<211> 198

<212> PRT

<213> Homo sapiens

<400> 175

Met Pro Leu Ala Gln Leu Ala Asp Pro Trp Gln Lys Met Ala Val Glu
 1 5 10 15

Ser Pro Ser Asp Ser Ala Glu Asn Gly Gln Gln Ile Met Asp Glu Pro
 20 25 30

Met Gly Glu Glu Glu Ile Asn Pro Gln Thr Glu Glu Val Ser Ile Lys
 35 40 45

Glu Ile Ala Ile Thr His His Val Lys Glu Gly His Glu Lys Ala Asp
 50 55 60

Pro Ser Gln Phe Glu Leu Leu Lys Val Leu Gly Gln Gly Ser Phe Gly
 65 70 75 80

Lys Val Phe Leu Val Lys Lys Ile Ser Gly Ser Asp Ala Arg Gln Leu
 85 90 95

Tyr Ala Met Lys Val Leu Lys Lys Ala Thr Leu Lys Val Arg Asp Arg
 100 105 110

Val Arg Thr Lys Met Glu Arg Asp Ile Leu Val Glu Val Asn His Pro
 115 120 125

Phe Ile Val Lys Leu His Tyr Ala Phe Gln Thr Glu Gly Lys Leu Tyr
 130 135 140

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Leu Ile Leu Asp Phe Leu Arg Gly Gly Asp Leu Phe Thr Arg Leu Ser
145 150 155 160

Lys Glu Val Met Phe Thr Glu Glu Asp Val Lys Phe Tyr Leu Ala Glu
165 170 175

Leu Ala Leu Ala Leu Asp His Leu His Ser Leu Gly Ile Ile Tyr Arg
180 185 190

Asp Leu Lys Pro Glu Lys
195

<210> 176

<211> 489

<212> PRT

<213> Homo sapiens

<400> 176

Met Ser Thr Glu Ala Asp Glu Gly Ile Thr Phe Ser Val Pro Pro Phe
1 5 10 15

Ala Pro Ser Gly Phe Cys Thr Ile Pro Glu Gly Gly Ile Cys Arg Arg
20 25 30

Gly Gly Ala Ala Ala Val Gly Glu Gly Glu Glu His Gln Leu Pro Pro
35 40 45

Pro Pro Pro Gly Ser Phe Trp Asn Val Glu Ser Ala Ala Ala Pro Gly
50 55 60

Ile Gly Cys Pro Ala Ala Thr Ser Ser Ser Ser Ala Thr Arg Gly Arg
65 70 75 80

Gly Ser Ser Val Gly Gly Gly Ser Arg Arg Thr Thr Val Ala Tyr Val
85 90 95

Ile Asn Glu Ala Ser Gln Gly Gln Leu Val Val Ala Glu Ser Glu Ala
100 105 110

Leu Gln Ser Leu Arg Glu Ala Cys Glu Thr Val Gly Ala Thr Leu Glu
115 120 125

Thr Leu His Phe Gly Lys Leu Asp Phe Gly Glu Thr Thr Val Leu Asp
130 135 140

Arg Phe Tyr Asn Ala Asp Ile Ala Val Val Glu Met Ser Asp Ala Phe
145 150 155 160

Arg Gln Pro Ser Leu Phe Tyr His Leu Gly Val Arg Glu Ser Phe Ser
165 170 175

Met Ala Asn Asn Ile Ile Leu Tyr Cys Asp Thr Asn Ser Asp Ser Leu
180 185 190

Gln Ser Leu Lys Glu Ile Ile Cys Gln Lys Asn Thr Met Cys Thr Gly
195 200 205

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[illegible]

<210> 177

<211> 105

<212> PRT
<213> Homo sapiens

<400> 177

Met Arg Glu Phe Glu Val Leu Lys Lys Leu Asn His Lys Asn Ile Val
1 5 10 15

Lys Leu Phe Ala Ile Glu Glu Glu Thr Thr Thr Arg His Lys Val Leu
20 25 30

Ile Met Glu Phe Cys Pro Cys Gly Ser Leu Tyr Thr Val Leu Glu Glu
35 40 45

Pro Ser Asn Ala Tyr Gly Leu Pro Glu Ser Glu Phe Leu Ile Val Leu
50 55 60

Arg Asp Val Val Gly Gly Met Asn His Leu Arg Glu Asn Gly Ile Val
65 70 75 80

His Arg Asp Ile Lys Pro Gly Asn Ile Met Arg Ala Leu Tyr His Ser
85 90 95

Leu Val Asp Asp Ser Phe His Pro Pro
100 105

<210> 178

<211> 413

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(413)

<223> "XAA" can be any amino acid

<400> 178

Met Tyr Cys Phe Gly Arg Lys Xaa Tyr Ile Ser Thr Arg Pro Cys Phe
1 5 10 15

Pro Asn Lys Thr Cys Gln Lys Met Leu Ile Ile Leu Thr Ser Ala Leu
20 25 30

Gln Ile Ala His Arg Cys Ile Cys Arg Ile Leu Leu Gly Ser Arg Val
35 40 45

Leu Ala Ala Lys Ala Ser Gly Asn Cys Thr Leu Asn Ser Glu Asp Phe
50 55 60

Ile Phe Asn Ile Gly Ser Ala Ala Tyr Asp Ala Val Leu Asp Arg Asn
65 70 75 80

Val Ala Ile Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln Thr His Ala
85 90 95

Lys Arg Ala Tyr Arg Glu Leu Val Leu Met Lys Cys Val Asn His Lys
100 105 110

Asn Ile Ile Ser Leu Leu Asn Val Phe Thr Pro Gln Lys Thr Leu Glu
115 120 125

Glu Phe Gln Asp Val Tyr Leu Val Met Glu Leu Met Asp Ala Asn Leu
130 135 140

Cys Gln Val Ile Gln Met Glu Leu Asp His Glu Arg Met Ser Tyr Leu
145 150 155 160

Leu Tyr Gln Met Leu Cys Gly Ile Lys His Leu His Ser Ala Gly Ile
165 170 175

Ile His Arg Asp Leu Lys Pro Ser Asn Ile Val Val Lys Ser Asp Cys
180 185 190

Thr Leu Lys Ile Leu Asp Phe Gly Leu Ala Arg Thr Ala Gly Thr Ser
195 200 205

Phe Met Met Thr Pro Tyr Val Val Thr Arg Tyr Tyr Arg Ala Pro Glu
210 215 220

Val Ile Leu Gly Met Gly Tyr Lys Glu Asn Val Asp Ile Trp Ser Val
225 230 235 240

Gly Cys Ile Met Gly Glu Met Val Arg His Lys Ile Leu Phe Pro Gly
245 250 255

Arg Asp Tyr Ile Asp Gln Trp Asn Lys Val Ile Glu Gln Leu Gly Thr
260 265 270

Pro Cys Pro Glu Phe Met Lys Lys Leu Gln Pro Thr Val Arg Asn Tyr
275 280 285

Val Glu Asn Arg Pro Lys Tyr Ala Gly Leu Thr Phe Pro Lys Leu Phe
290 295 300

Pro Asp Ser Leu Phe Pro Ala Asp Ser Glu His Asn Lys Leu Lys Ala
305 310 315 320

Ser Gln Ala Arg Asp Leu Leu Ser Lys Met Leu Val Ile Asp Pro Ala
325 330 335

Lys Arg Ile Ser Val Asp Asp Ala Leu Gln His Pro Tyr Ile Asn Val
340 345 350

Trp Tyr Asp Pro Ala Glu Val Glu Ala Pro Pro Pro Gln Ile Tyr Asp
355 360 365

Lys Gln Leu Asp Glu Arg Glu His Thr Ile Glu Glu Trp Lys Glu Leu
370 375 380

Ile Tyr Lys Glu Val Met Asn Ser Glu Glu Lys Thr Lys Asn Gly Val
385 390 395 400

Val Lys Gly Gln Pro Ser Pro Ser Ala Gln Val Gln Gln
405 410

<210> 179
<211> 108
<212> PRT
<213> Homo sapiens

<400> 179

Met Ser Lys Ser Lys Val Asp Asn Gln Phe Tyr Ser Val Glu Val Gly
1 5 10 15

Asp Ser Thr Phe Thr Val Leu Lys Arg Tyr Gln Asn Leu Lys Pro Ile
20 25 30

Gly Ser Gly Ala Gln Gly Ile Val Cys Ala Ala Tyr Asp Ala Val Leu
35 40 45

Asp Arg Asn Val Ala Ile Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln
50 55 60

Thr His Ala Lys Arg Ala Tyr Arg Glu Leu Val Leu Met Lys Cys Val
65 70 75 80

Asn His Lys Asn Val Ser Phe Val Ile Phe Lys Leu Leu Ala Val Gly
85 90 95

Val Cys Lys Ile Gly Lys Arg Lys Cys Val Cys Thr
100 105

<210> 180
<211> 336
<212> PRT
<213> Homo sapiens

<400> 180

Met Ala Met Thr Gly Ser Thr Pro Cys Ser Ser Met Ser Asn His Thr
1 5 10 15

Lys Glu Arg Val Thr Met Thr Lys Val Thr Leu Glu Asn Phe Tyr Ser
20 25 30

Asn Leu Ile Ala Gln His Glu Glu Arg Glu Met Arg Gln Lys Lys Leu
35 40 45

Glu Lys Val Met Glu Glu Glu Gly Leu Lys Asp Glu Glu Lys Arg Leu
50 55 60

Arg Arg Ser Ala His Ala Arg Lys Glu Thr Glu Phe Leu Arg Leu Lys
65 70 75 80

Arg Thr Arg Leu Gly Leu Glu Asp Phe Glu Ser Leu Lys Val Ile Gly
85 90 95

Arg Gly Ala Phe Gly Glu Val Arg Leu Val Gln Lys Lys Asp Thr Gly

100

105

110

His Val Tyr Ala Met Lys Ile Leu Arg Lys Ala Asp Met Leu Glu Lys
 115 120 125

Glu Gln Val Gly His Ile Arg Ala Glu Arg Asp Ile Leu Val Glu Ala
 130 135 140

Asp Ser Leu Trp Val Val Lys Met Phe Tyr Ser Phe Gln Asp Lys Leu
 145 150 155 160

Asn Leu Tyr Leu Ile Met Glu Phe Leu Pro Gly Gly Asp Met Met Thr
 165 170 175

Leu Leu Met Lys Lys Asp Thr Leu Thr Glu Glu Glu Thr Gln Phe Tyr
 180 185 190

Ile Ala Glu Thr Val Leu Ala Ile Asp Ser Ile His Gln Leu Gly Phe
 195 200 205

Ile His Arg Asp Ile Lys Pro Asp Asn Leu Leu Leu Asp Ser Lys Gly
 210 215 220

His Val Lys Leu Ser Asp Phe Gly Leu Cys Thr Gly Leu Lys Lys Ala
 225 230 235 240

His Arg Thr Glu Phe Tyr Arg Asn Leu Asn His Ser Leu Pro Ser Asp
 245 250 255

Phe Thr Phe Gln Asn Met Asn Ser Lys Arg Lys Ala Glu Thr Trp Lys
 260 265 270

Arg Asn Arg Arg Gln Leu Ala Phe Ser Thr Val Gly Thr Pro Asp Tyr
 275 280 285

Ile Ala Pro Glu Val Phe Met Gln Thr Gly Tyr Asn Lys Leu Cys Asp
 290 295 300

Trp Trp Ser Leu Gly Val Ile Met Tyr Glu Met Leu Ile Gly Lys Leu
 305 310 315 320

His Gly Phe Arg Gly Leu Phe Leu Cys Ile His Asp Arg Leu Leu His
 325 330 335

<210> 181

<211> 415

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(415)

<223> "XAA " can be any amino acid

<400> 181

Xaa Arg His Glu Ser Ala Arg Ala Ala Arg Val Ser Gly Gly Ser Met
1 5 10 15

Leu Asp Ile Ile Lys Tyr Ile Val Asn Arg Gly Glu His Lys Asn Gly
20 25 30

Val Leu Glu Glu Ala Ile Ile Ala Thr Ile Leu Lys Glu Val Leu Glu
35 40 45

Gly Leu Asp Tyr Leu His Arg Asn Gly Gln Ile His Arg Asp Leu Lys
50 55 60

Ala Gly Asn Ile Leu Leu Gly Glu Asp Gly Ser Val Gln Ile Ala Asp
65 70 75 80

Phe Gly Val Ser Ala Phe Leu Ala Thr Gly Gly Asp Val Thr Arg Asn
85 90 95

Lys Val Arg Lys Thr Phe Val Gly Thr Pro Cys Trp Met Ala Pro Glu
100 105 110

Val Met Glu Gln Val Arg Gly Tyr Asp Phe Lys Ala Asp Met Trp Ser
115 120 125

Phe Gly Ile Thr Ala Ile Glu Leu Ala Thr Gly Ala Ala Pro Tyr His
130 135 140

Lys Tyr Pro Pro Met Lys Val Leu Met Leu Thr Leu Gln Asn Asp Pro
145 150 155 160

Pro Thr Leu Glu Thr Gly Val Glu Asp Lys Glu Met Met Lys Lys Tyr
165 170 175

Gly Lys Ser Phe Arg Lys Leu Leu Ser Leu Cys Leu Gln Lys Asp Pro
180 185 190

Ser Lys Arg Pro Thr Ala Ala Glu Leu Leu Lys Cys Lys Phe Phe Gln
195 200 205

Lys Ala Lys Asn Arg Glu Tyr Leu Ile Glu Lys Leu Leu Thr Arg Thr
210 215 220

Pro Asp Ile Ala Gln Arg Ala Lys Lys Val Arg Arg Val Pro Gly Ser
225 230 235 240

Ser Gly His Leu His Lys Thr Glu Asp Gly Asp Trp Glu Trp Ser Asp
245 250 255

Asp Glu Met Asp Glu Lys Ser Glu Glu Gly Lys Ala Ala Phe Ser Gln
260 265 270

Glu Lys Ser Arg Arg Val Lys Glu Glu Asn Pro Glu Ile Ala Val Ser
275 280 285

Ala Ser Thr Ile Pro Glu Gln Ile Gln Ser Leu Ser Val His Asp Ser
290 295 300

Gln Gly Pro Pro Asn Ala Asn Glu Asp Tyr Arg Glu Ala Ser Ser Cys
305 310 315 320

Ala Val Asn Leu Val Leu Arg Leu Arg Asn Ser Arg Lys Glu Leu Asn
325 330 335

Asp Ile Arg Phe Glu Phe Thr Pro Gly Arg Asp Thr Ala Asp Gly Val
340 345 350

Ser Gln Glu Leu Phe Ser Ala Gly Leu Val Asp Gly His Asp Val Val
355 360 365

Ile Val Ala Ala Asn Leu Gln Lys Ile Val Asp Asp Pro Lys Ala Leu
370 375 380

Lys Thr Leu Thr Phe Lys Leu Ala Ser Gly Cys Asp Gly Ser Glu Ile
385 390 395 400

Pro Asp Glu Val Lys Leu Ile Gly Phe Ala Gln Leu Ser Val Ser
405 410 415

<210> 182
<211> 409
<212> PRT
<213> Homo sapiens

<220>
<221> -
<222> (1)..(409)
<223> "Xaa" can be any amino acid

<400> 182

Xaa Arg His Glu Ser Ala Arg Ala Ala Arg Val Ser Gly Gly Ser Met
1 5 10 15

Leu Asp Ile Ile Lys Tyr Ile Val Asn Arg Gly Glu His Lys Asn Gly
20 25 30

Val Leu Glu Glu Ala Ile Ile Ala Thr Ile Leu Lys Glu Val Leu Glu
35 40 45

Gly Leu Asp Tyr Leu His Arg Asn Gly Gln Ile His Arg Asp Leu Lys
50 55 60

Ala Gly Asn Ile Leu Leu Gly Glu Asp Gly Ser Val Gln Ile Ala Asp
65 70 75 80

Phe Gly Val Ser Ala Phe Leu Ala Thr Gly Gly Asp Val Thr Arg Asn
85 90 95

Lys Val Arg Lys Thr Phe Val Gly Thr Pro Cys Trp Met Ala Pro Glu
100 105 110

Val Met Glu Gln Val Arg Gly Tyr Asp Phe Lys Ala Asp Met Trp Ser
115 120 125

